

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number 1

TO: Andrew D Kosar Location: 3c04 / 3c18

Art Unit: 1654

Wednesday, May 10, 2006

Case Serial Number: 10/800179

From: Noble Jarrell

Location: Biotech-Chem Library

Rem 1B71

Phone: 272-2556

Noble.jarrell@uspto.gov

Search Notes	



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STIC-Biotech/ChemLib

From: Sent: To:

Subject:

189439

Requester: ANDREW KOSAR (P/1654) Art Unit: GROUP ART UNIT 1654 Employee Number: 80341 Office Location: REM 03C04 Phone Number: (571)272-0913 Mailbox Number: REM 3c18		
Case serial number:		
10/800,179 Class / Subclass(es):		
Earliest Priority Filing Date	: :	
Format preferred for results: Paper Search Topic Information: Please search SEQ ID NO:		
Special Instructions and Othe	er Comments:	
***************** Searcher: Searcher Phone: Date Searcher Picked up: Date completed: Searcher Prep Time: Online Time:	************ Type of Search NA# AA#: S/L: Oligomer: Encode/Transl: Structure #: Text: Inventor: Litigation:	Vendors and cost where applicable STN: DIALOG: QUESTEL/ORBIT: LEXIS/NEXIS:_ SEQUENCE SYSTEM: Compley WWW/Internet: Other (Specify):

ANDREW KOSAR [andrew.kosar@uspto.gov] Tuesday, May 09, 2006 5:16 PM STIC-Biotech/ChemLib

Database Search Request, Serial Number: 10/800,179

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PALM INTRANET

Day: Wednesday Date: 5/10/2006 Time: 16:58:59

Inventor Information for 10/800179

Inventor Name	City	State/Country
KUMAR, MANOJ	FREMONT	CALIFORNIA
CUEVAS, WILLIAM A.	SAN FRANCISCO	CALIFORNIA
Appln Info Contents Petition Info	o Atty/Agent Info Contir	nuity Data Foreign Data In
7 Calley III		nuity Data Foreign Data In
	Search or Patent#	
Search Another: Application# PCT //		# Search
Search Another: Application#	Search or Patent#	# Search Search

To go back use Back button on your browser toolbar.

Back to PALM | ASSIGNMENT | OASIS | Home page

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Protein Sequence Searches - February 2005

All of the sequence databases on ABSS have recently been updated.

- Please note that the curators of the UniProt database have purged some temporary accession numbers from the most recent version of UniProt. These sequences have been assigned new permanent accession numbers. The new UniProt record may not contain the previous temporary accession number.
- If you encounter an accession number from an older search run against UniProt (results file extension .rup) that can no longer be found in the database, the permanent record with the new accession number can be found by searching the old accession number in the UniProt Protein Archive database (UniPARC) at:

http://www.pir.uniprot.org/database/archive.shtml

If you have any questions regarding this information or your results, please contact any STIC searcher.

When submitting sequence search results for scanning into IFW, please include a copy of this attachment to assist any future Examiners or members of the public who may encounter UniProt temporary accession numbers.

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November 2005

Published_Applications Nucleic Acid and Published_Applications Amino Acid database searches now generate two sets of results each. The Published_Applications databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Newly published applications will appear in the Published_Applications_New databases; older published applications make up the Published_Applications_Main databases.

Searches run against Nucleic Acid Published_Applications produce two sets of results, with the extensions .rnpbm (Published_Applications_NA_Main) and .rnpbn (Published_Applications_NA_New).

Searches run against Amino Acid Published_Applications produce two sets of results, with the extensions .rapbm (Published_Applications_AA_Main) and .rapbn (Published_Applications_AA_New).

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Result
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Maximum DB
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ALIGNMENTS

R;Bressan, G.M.; Argos, P.; Stanley, K.K. Biochemistry 26, 1497-1503, 1987 A;Title: Repeating structure of chick tropoelastin revealed by complementary DNA cloning A;Reference number: A26601; MUID:87242320; PMID:3593675

;Species: Gallus gallus (chicken) ;Date: 05-Oct-1988 #sequence_revision 26-Jul-1996 #text_change 09-Jul-2004 ;Accession: A26601; A30795; A27264

A; Accession: A26601

N; Alternate names: tropoelastin

lastin precursor - chicken (fragment)

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A; Molecule type: mRNA
A; Molecule type: mRNA
A; Residues: 491-569, 'G',571-604,'A',606-643,'A',645-687,'R',689-700,'R',702-784 <TOK>
A; Residues: 491-569, 'G',571-604,'A',606-643,'A',645-687,'R',689-700,'R',702-784 <TOK>
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C; Superfamily: elastin
C; Keywords: alternative splicing; extracellular matrix; glycoprotein; hydroxylysine
C; Keywords: alternative splicing; extracellular matrix; glycoprotein; hydroxylysine
F;1-24/Domain: signal sequence (fragment) #status predicted <NAT>
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A;Residues: 1-212,237-524,535-784 <BR2>
A;Residues: 1-212,237-524,535-784 <BR2>
A;Cross-references: UNIPROT:P07916; UNIPARC:UPI0000129E6D; GB:M15889; NID:g212803; PIDN:#
R;Baule, V.J.; Foster, J.A.
Biochem. Biophys. Res. Commun. 154, 1054-1060, 1988
A;Title: Multiple chick tropoelastin mRNAs.
A;Title: Multiple chick tropoelastin mRNAs.
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A;Residues: 85-784 cBAU>
A;Residues: 85-784 cBAU>
A;Cross-references: UNIPARC:UDI00001713B4; GB:M21880; NID:g212741; PIDN:AAA49082.1; PID:gR;Tokimitsu, I.; Tajima, S.; Nishikawa, T.; Tajima, M.; Fukasawa, T.
A;Tokimitsu, I.; Tajima, S.; Nishikawa, T.; Tajima, M.; Fukasawa, T.
Arch. Biochem. Biophys. 256, 455-461, 1987
A;Title: Sequence analysis of elastin cDNA from chick aorta and tissue-specific transcrif A;Reference number: A27264; MUID:87297534; PMID:3502711
A;Recession: A27264
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AAKAGAGLGGVGGIGGLGGVGGVGVPGGLGVPGVVQPGVGAAGKPPKVPGAGIPGAFPGG
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37; Mismatches 188;
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elastin precursor - mouse

N,Alternate names: tropoelastin
C;Species: Mus musculus (house mouse)
C;Date: 18-Aug-1995 #sequence_revision 16-Aug-1996 #text_chan
C;Accession: A55721
R;Wydner, K.S.; Sechler, J.L.; Boyd, C.D.; Passmore, H.C.
Genomics 23, 125-131, 1994
A;Title: Use of an intron length polymorphism to localize the
A;Reference number: A55721
A;Accession: A55721
A;Accession: A55721
A;Molecule type: mRNA
A;Residues: 1-860 <WYD>
A;Cross-references: UNIPROT:P54320; UNIPARC:UPI0000029BE6; GB
C;Genetics:
A;Cross-references: UNIPROT:P54320; UNIPARC:UPI0000029BE6; GB
C;Superfamily: elastin
C;Keywords: alternative splicing; extracellular matrix; glyco
F;1-27/Domain: signal sequence #status predicted <MATI>
F;850-855/Disulfide bonds: #status predicted
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;Species: Mus musculus (house mouse)
;Date: 18-Aug-1995 #sequence_revision 16-Aug-1996 #text_change
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                                                                                                                                                                                                                                                   GSGAGAGSGAGAGSGAGAGSGV-GVPGVGVPGVGVP-GKGVPGVGPGVGPG--VGPGVGP
                                                                                                                                                                                                                                                                                                                                          GKPGK-VPGVGLPGVYPGGVLPGTGARFPGVGVLPGVPTGTGVKAKAPGGGGAFSGIFG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VGVPGKGVPGVG-PGVGP-GVGPGVG---PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPG
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                                              AGGLGAG-
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                                                                                                                                                                                                         G--AGAGSGAGAGSGAGAGS----GAGAGSGVGVPGVGVPGVGVP---
                                                                                                                                                                                                                               PAAAAAAAAKAAAKAGLGPGVGGVPG-GVGVGGIPGGVGVGGVGGVGPGGVTGIGAGP
                                                                                                                                                                                                                                                                             AKAAKYGAG---GAGALGGL-----
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                                                                                                                                                                                                                                                                                                                          PAAAAKAAKAAKYGARGGVGIPTYGV-GAG----GFPGYGVGAGAGLG-GASPAAAAAA
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                                                                                         VPGRVAGAAP-
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                                             -GLGAG
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Pred. No. 1.2e-66;
0; Mismatches 274
                                             -GLGAG
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                                             GLGAGGGVSPAAAAKAAKYGAAGLG
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RESULT 3

EART
elastin precursor - rat
N;Alternate names: tropoelastin
C;Species: Rattus norvegicus (Norway rat)
C;Date: 11-Jan-1991 #sequence revision 16-Aug-1996 #text_change 09-Jul-2004
C;Accession: A36106; A30878; \bar{A}36523; S02173; I54172; I68\bar{5}05
R;Pierce, R.A.; Deak, S.B.; Stolle, C.A.; Boyd, C.D.

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A;Residues; 22-31 <PRAN-
A;Residues; 27-31, MUID:89117149; PMID:2913947
A;Residues: (TP., 369-545, 548-764,770-864 <RIC>
A;Residues: (TP., 369-545, 548-764,770-864 <RIC)
A;Residues: (TP., 369-545, 548-764,770-864 <RIC)
A;Residues: (TP., 369-546,782)
A;Residues: (T
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A;Title: Heterogeneity of rat tropoelastin mRNA revealed by cDNA cloning.

A;Reference number: A36106; MUID:91104868; PMID:1702999

A;Recession: A36106

A;Molecule type: mRNA

A;Residues: 1-864 <PIE>
A;Cross-references: UNIPROT:Q99372; UNIPARC:UPI0000129E6F; GB:M60647; GB:J05292; NID:g20

A;Title: Rat tropoelastin is synthesized from a 3.5-kilobase mRNA.

A;Reference number: A30878; MUID:88330868; PMID:2971041

A;Reference number: A30878; MUID:88330868; PMID:2971041

A;Reterence number: A30878; MUID:89359327; PMID:2768256

A;Rolecule type: gracein

A;Residues: 22-31 craps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    h 28.4%; Score 1185; DB 1;
Similarity 43.8%; Pred. No. 1.1e-64;
78; Conservative 47; Mismatches 299;
                                                                                 GPGA-GAGSGAGAGSGAGAGSGAGAGSGVGVP--GVGVPGVGVPGKGVPGVG-PGVGP-G 161
                                                                                                                                                                                                                                                                                                                                                GVPGAVPGGVPG-GLPG-GVPGGVYYPGAGIGGGLGGGALGPGGKPPKPGAGLLGAFGAG
                                                                                                                                                                                                                                                                                                                                                                                                                             GVPGV---GVPGVGVPGKGVPG----VGPGVGPGVGPG-VGPGA----GAG----SGAG
VPGAVGVGGVPGAVGGIGGIGGLGVSTGAVVPQLGAGVGAGGKPGK-VPGVGLPGVYPGG
                                                                                                                                                                   PGGLGGAGPGAGLSYASRPGGVLVPGGGAGAAAAYKAAAKAGAGLGGIG-GVPGGVGVGG 139
                                                                                                                                                                                                                                                     AGSGAGAGSGAGAGSGVGVPGVGVPGVGV------PGKGVPGVGPGVGPGVGPGV 106
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198
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J. Biol. Chem. 262, 5/55-5/64, 1797
A;Title: Primary structures of bovine elastin a, b, and
A;Reference number: A92640; MUID:87194772; PMID:3032943
                                                                                                                                                                                                                               N;Alternate names: tropoelastin
N;Contains: elastin precursor, splice form b; elastin precursor,
C;Species: Bos primigenius taurus (cattle)
C;Date: 08-Uun-1989 #sequence revision 26-Jul-1996 #text change (C;Accession: A31865; A26728; B26728; C26728; A22343; I45886
R;Yeh, H.; Anderson, N.; Ornstein-Goldstein, N.; Bashir, M.M.; Re
                                                                                                                                                                         Biochemistry 28, 2365-2370, 1989
A; Title: Structure of the bovine elastin gene and S1 nuclease A; Reference number: A31865; MUID:89274159; PMID:2543440
                                                  A;Cross-references: UNIPROT:P04985; UNIPROT:Q28101; UNIPARC:UPI000016C304; GB:J02855; R;Raju, K.; Anwar, R.A.
J. Biol. Chem. 262, 5755-5762, 1987
                                                                                                                                                                                                                                                                                                                                                EABO elastin precursor, splice form a - bovine
                                                                                                                A; Molecule type: DNA
A; Residues: 1-27 < YEH>
                                                                                                                                                      A; Accession: A31865
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A;Molecule type: mRNA A;ReBidues: 1,'RS',4-11,'E',13-636,'V',638-747 <RAJ> A;Cross-references: UNIPARC:UPI0000129E6C; GB:J02717;

NID:g163019; PIDN:AAA30503.1;

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A; Molecule type: DNA
A; Molecule type: DNA
A; Residues: 678-683,685-747 < ROS>
A; Residues: 678-683,685-747 < ROS>
A; Cross-references: UNIPARC: UPI000008856A; GB:M31898; NID:g163015; PIDN:AAA96417.1;
A; Cross-references: UNIPARC: UPI000008856A; GB:M31898; NID:g163015; PIDN:AAA96417.1;
R; Brown, P.L.; Mecham, L.; Tisdale, C.; Mecham, R.P.
Blochem. Biophys. Res. Commun. 186, 549-555, 1992
A; Title: The cysteine residues in the carboxy terminal domain of tropoelastin form a A; Reference number: A58621; MUID:92337651; PMID:1632791
A; Contents: annotation, disulfide bonds
C; Comment: The term tropoelastin refers to a soluble precursor form of the extracell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Note: the list of introns is incomplete
C;Superfamily: elastin
C;Keywords: alternative splicing; extracellular matrix; glycoprotein; hydroxylysine
F;1-747/Product: elastin precursor, splice form a #status predicted <EPA>
F;1-225,260-747/Product: elastin precursor, splice form c #status predicted <EPC>
F;1-225,240-747/Product: elastin precursor, splice form b #status predicted <EPB>
F;1-26/Domain: signal sequence #status predicted <SIG>
F;1-26/Domain: signal sequence #status predicted <ANT>
F;27-747/Product: elastin #status predicted <MAT>
F;105,109,252,271,275,324,327,400,404,407,445,448,489,493,544,548,552,606,609,645,64
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A;Accession: A22343
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A;Cross-references: UNIDARC:UPIO00002A8D4; GB:K03566; NID:g1C
R;Cicila, G.; May, M.; Ornstein-Goldstein, N.; Indik, Z.; Moi Biochemistry 24, 3075-3080, 1985
A;Title: Structure of the 3, portion of the bovine elastin go
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A;Accession: Cype: mRNA
A;Residues: 1,'RS',4-11,'E',13-225,240-636,'V',638-747 <RA2>
A;Cross-references: UNIPARC:UPI000002A8D3; GB:K03505; NID:g163025; PIDN:AAA30505.1;
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A;Title: Biology of disease: Elastin: Relation of protein
A;Reference number: I45885; MUID:85059254; PMID:6150137
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A; Residues: 613-747 < CIC>
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                                                                             LGAGVGAGVKPGK-----VPGVGLPGV-YPGGVLPGAGARFPGI--GVLPGV
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                              - APGAI PGI GGI AGVGAPDAAAAAAAAAKAAKFGAAGGL PGVGVPGVGVP
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41.4%; Pred. No. 1.1e-62;
tive 46; Mismatches 249)
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RESULT 5

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troppelastin - sheep
C;Species: Ovis orientalis aries, Ovis ammon aries (don C;Species: Ovis orientalis aries, Ovis ammon aries (don C;Date: 23-Aug-1996 #sequence_revision 13-Mar-1997 #tes
C;Accession: S59623; A24758
R;Mauch, J.C.; Sandberg, L.B.; Roos, P.J.; Jimenez, F.;
Rauch, J.C.; Sandberg, L.B.; Roos, P.J.; Jimenez, F.;
Matrix Biol. 14, 635-641, 1994
A;Title: Extensive alternate exon usage at the 5' end of A;Reference number: S59623
A;Accession: S59623
A;Accession: S59623
A,Status: preliminary; not compared with conceptual transcriptors.
                                                                                                                                                  A;Cross-references: UNIPROT:P11547; UNIPARC:UPI0000177409
R;Yoon, K.; Davidson, J.M.; Boyd, C.; May, M.; LuValle, P.; Ornstein-Goldstein, N.;
Arch. Biochem. Biophys. 241, 684-691, 1985
A;Title: Analysis of the 3' region of the sheep elastin gene.
A;Reference number: A24758; MUID:85305763; PMID:3839997
A;Accession: A24758
A;Rolecule type: mRNA
A;Residues: 655-669,671-716,732-770 <YOO>
A;Cross-references: UNIPARC:UPI0000129E70
C;Superfamily: elastin
C;Reywords: alternative splicing; extracellular matrix; glycoprotein; hydroxylysine
F;760-765/Disulfide bonds: #status predicted
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A; Residues: 1-770 < MAU>
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                                         GSGAGAGSGVGVP-GVGVPGVGVPGKGVPGVGPGV---GPGVGPGVGPGAGA-GSGA---
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   GGVLGAVPG-GVPGGVFFPGAGLGGLGVGGLGPGVKPAKPGVGGLVGPGLGAEGSGLLPA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  --AAAAAAKAAAKAAQFGLG-GVGGLGVGGLGAVPG----AVGLG---GVSPAAAAKAAK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -----AAKAAKIGAGGVGALGGVVP--GAPGAIPGL-PGVGGVPGVGIPAAAAAKAAA
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                                                                             Conservative
                                                                                                 42.5%;
                                                                                                                27.5%;
                                                                               41; Mismatches
                                                                                               Score 1146.5; DB Pred. No. 2.1e-62;
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#text change 09-Jul-2004
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Smith

В

57

-AGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPG 101

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period clock protein - mouse (fragment)
(;Species: Mus musculus (house mouse)
(;Date: 31-Mar-1988 #sequence_revision 31-Mar-1988 #text_change 09-Jul-2004
(;Accession: A24403
R;Shin, H.S.; Bargiello, T.A.; Clark, B.T.; Jackson, F.R.; Young, M.W.
Nature 317, 445-448, 1985
A;Title: An unusual coding sequence from a Drosophila clock gene is conserve
A;Reference number: A24403; MUID:86014384; PMID:2413365
A;Accession: A24403
A;Molecule type: DNA
A;Residues: 1-713 <SHI>
A;Residues: 1-713 <SHI>
A;Cross-references: UNIPROT:P08399; UNIPARC:UPI000016D179; GB:X02966; GB:M12039; NID:955
C;Comment: Mutations within the per locus of the fruit fly affect a variety of natural h
logous locus with multiple tandem repeats of nucleic acid hexamers (ACNGGN, TCAGGC) that
C;Comment: The serine residues of the S-G repeats found in certain proteoglycans are att
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UMMS
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                                                                                                                                                                                                                                                                                                       PFGGALGALGFPG---
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Matches
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;Keywords: circadian rhythm; tandem
;41-77/Domain: EGF homology <EGF>
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nilarity 35.3%;
Conservative 54
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RESULT 7

F70806

hypothetical glycine-rich protein Rv3508 - Mycobacterium tuberculosis (strain C; Species: Mycobacterium tuberculosis
C; Species: Mycobacterium tuberculosis
C; Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 09-Jul-2004
C; Accession: F70806
R; Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D. R; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, Kajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, Nature 393, 537-544, 1998
Nature 393, 537-544, 1998
A; Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.C.
A; Title: Deciphering the biology of Mycobacterium tuberculosis from the complex of the

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A;Accession: F70806
A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
A;Residues: 1-1901 <COL>
A;CCOLS
A;CCOSS-references: UNIPROT:053553; UNIPARC:UPI000013C2A6; GB:AL022022; GB:AL123456; NID
A;Experimental source: strain H37Rv
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C;Superfamily:
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1; Mismatches 368;
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A;Accession: B70807
A;Status: preliminary; nucleic acid sequence not shown; translation not sh A;Molecule type: DNA
A;Residues; 1-1079 <CCL>
A;Cross-references: UNIPARC:UPI0000D3AEA; GB:AL022022; GB:AL123456; NID:9A;Cross-references: UNIPARC:UPI0000D3AEA; GB:AL022022; GB:AL123456; NID:9A;Status: R73512
C;Superfamily: collagen alpha 1(I) chain; fibrillar collagen carboxyl-term
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Matches 317; Conserv
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Conservative

26.1%; Sc 36.6%; Pr tive 32;

Score 1089.5; DB 2; Pred. No. 6.9e-59; 2; Mismatches 394;

Indels Length

123; 1079

Gaps

31;

fibrillar collagen carboxyl-terminal homology;

NID: g3261554;

complete B . G

genome

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Holroyd,

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Gordon,

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1274 GGDGGNAGFGAGVPGDGGDGGNAGFGAGVPGDGGIGGTGGAGGAGGAGA 1322
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VPGVGVPGKGVPGVGPG : GAGIGSLGGGTGGDGGN	VGPGAGAGSGAGA- FGGGAGAGGGGLTAGAN		GSGVGVPGVGVPGVGVP : : : GNGGIAGMG	gsgygypgygyp 		GSGVGVPGVGVP ; AGGTGGNGGNITGGTAG				
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R; Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Gor; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Holro Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S. Nature 393, 537-544, 1998

A; Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G. A; Title: Deciphering the biology of Mycobacterium tuberculosis from the complete g. R; Reference number: A70500; MUID:98295987; PMID:9634230

A; Reference number: A70500; MUID:98295987; PMID:9634230

A; Status: preliminary; nucleic acid sequence not shown; translation not shown A; Molecule type: DNA

A; Residues: 1-1489 cOLL>
A; Cross-references: UNIPARC:UPI00000D3AEB; GB:AL022022; GB:AL123456; NID:93261554; A; Gene: Rv3514

C; Superfamily: collagen alpha 1(IV) chain
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C;Species: Mycobacterium tuberculosis
C;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 20-Jun-2000
C;Accession: D70807
C;Accession: D70807
R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.;
R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGS
                                                                                                                                                                                                                                                                                                                               VGVPGVGVPG---VGVPGK-GVPGVGPGVGPGVGPGVG---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGGAAGTGGTGGMIGTTGNAGVGGAG---GQGGDGGAGGAGADADQPGATGGTGFAGGAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GAGAGSGAGAGSGVGVPG-VGVPGVGVFGKGVPGVGPGVG-----PGVGPGVGPGAGAG-
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  GGTGGVGGTGGDGGNAGTGAGDPGKG-GTGGTGGTGGSGGAGGSGGANFNGGTGGTGGTG
                                                                                                                                                                                     GIGGTGGDGGTGGAAGAGGAAGTGGTGGMIGTTGNAGVGGAGGQGGDGGAGGAGADA
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                                                                                                                                         PGAGAGSGAGAGSGAGAGSG
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                                              -GVPGKGVPGVGPGVG---PGVGPGVGPGAGAGSGAGAGSGAGA----GSGAGAGSG 313
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36.0%;
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                                                                                                                                         AGAGSGVGVPGVGVPGV-----
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                                                                                                                                                                                                                                                                                                       GAGAGSGAGAGSGAGAGSGVGVPGVGVPG--VGVPGKGVPGVGPGVGPGV--GP
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                               AGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGAGAGSGAGA
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 -GNGGNGGTGVGSPGGLGGAGGTGGLGGAGAGGGA
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A;Molecule type: mRNA
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-453,483-617,651-792 <FAZ>
A;Note: this sequence represents a composite of several splice forms
A;Residues: 1-453,483-617, Kuivaniemi, H.; Chu, M.L.; Davidson, J.M.; Rosenbloom, J.;
R;Fazio, M.J.; Olsen, D.R.; Kuivaniemi, H.; Chu, M.L.; Davidson, J.M.; Rosenbloom, J.;
Lab. Invest. 58, 270-277, 1988
A;Title: Isolation and characterization of human elastin cDNAs, and age-associated var
A;Reference number: A53891; MUID:88156138; PMID:2831431 A;Cross-references: UNIDARC:UPI000006E12E; GB:J04821; NID:g182052; R;Fazio, M.J.; Olsen, D.R.; Kauh, E.A.; Baldwin, C.T.; Indik, Z.; O J. Invest. Dermatol. 91, 458-464, 1988
A;Title: Cloning of full-length elastin cDNAs from a human skin fib A;Reference number: A30524; MUID:89009960; PMID:3171221
A;Accession: A30524 elastin precursor, long splice in RALTernate names: tropoclastin C; Species: Homo sapiens (man) C; Date: 22-Jun-1990 #sequence roc; Accession: A32707; A33705; A3 A;Title: Characterization of the complete human elastin A;Reference number: A33705; MUID:89255358; PMID:2722804 A;Accession: A33705 A; Molecule type: mRNA A; Residues: 1-500,507-792 <IND> A; Cross-references: UNIPROT: P15502; R;Indik, Z.; Yeh, H.; Ornstein-Goldstein, N.; Sheppard, P.; Ar Proc. Natl. Acad. Sci. U.S.A. 84, 580-5864, 1987 A;Title: Alternative splicing of human elastin mRNA indicated A;Reference number: A32707; MUID:87289668; PMID:3039501 R;Bashir, M.M.; Indik, Z.; Yeh, H.; J. Biol. Chem. 264, 8887-8891, 1989 A;Molecule type: DNA A;Residues: 1-27 <BAS> A; Accession: A32707 Species: Homo sapiens (man)

Date: 22-Jun-1990 #sequence revision 26-Jul-1996 #text_change 09-Jul-2004;

Accession: A32707; A33705; A30524; A53891 10 form Ornstein-Goldstein, UNIPROT:Q9UMK5; UNIPARC:UPI0000173C4E; GB:M16983; Ornstein-Goldstein, N.; Rosenbloom, J.C.; Abrams, gene. P.; Anderson, Delineation à sequence Ornstein-Goldstein, PIDN:AAA52379.1; PID: Ornstein-Goldstein, N. N.; Rosenbloom, analysis of cl of unusual

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A;Cross-references: GDB:119107; OMIM:130160
A;Map position: 7g11.23-7g11.23
C;Superfamily: elastin
C;Keywords: alternative splicing; extracellular matrix;
F;1-26/Domain: signal sequence #status predicted <NAT>
F;27-792/Product: elastin #status predicted <MAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Accession: A53891
A;Accession: A53891
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 164-453,483-500,507-617,651-792 <FA2>
A;Cross-references: UNIPARC:UPI000016A887; GB:M24782; NID:g182063; PIDN:AAA53190.1; PID:
A;Cross-references: UNIPARC:UPI000016A887; GB:M24782; NID:g182063; PIDN:AAA53190.1; PID:
C;Comment: The term tropoelastin refers to a soluble precursor form of the extracellular
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VGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGAGAGSGAGAG 779
                          VPGALAAAKAAKYGAAVPGVLGGLGALGGVGI-PGGVVGAGPAAAAAAAKAAAKAAQFGL
                                                     VPG-VGVPGVGVPGVGPGVGPGVGPGVGPGVGPGAGAGSGAGAGSGAGAGSGAGAGSG-
                                                                                                        PGKGVPGVGPGVGPGVGPGVG-PGAGAGSGAGA-----GSGAGAGSGAGAGSGVG
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                                                                                                                                                                                                                                                                     YGARPGVGVGGIPTYGVGAGGFPGFGVGVG-----GIPGVAGVPSVG-----GVPGVG-GV
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40.6%; Pred. No. 6e-5
tive 47; Mismatches
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Db 710 VGAAGLG--GLGVGGLGVPGVG-GLG-GIPPAAAAKAAKYGAAGLG 751
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R;Xu, M.; Lewis, R.V.
proc. Natl. Acad. Sci. U.S.A. 87, 7120-7124, 1990
A;Title: Structure of a protein superfiber: spider dragline
A;Reference number: A36068; MUID:90384959; PMID:2402494
A;Accession: A36068
A;Status: preliminary
A;Molecule type: DNA
A;Molecule type: DNA
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A;Residues: 1-718 <XUA>
A;Cross-references: UNIPROT:P19837; UNIPARC:UPI000016BF56; GB:M37137;
A;Note: the authors translated the codon GGT for residue 292 as Gln, C
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hypothetical glycine-rich protein Rv2490c - Mycobacterium tuberculosis (strain H37RV) C;Species: Mycobacterium tuberculosis (c;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 09-Jul-2004 C;Accession: A70869

R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Gordon, S.; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Holroyd, S.; Connor, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S. Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S. Nature 393, 537-544, 1998

A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G. A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete genome A;Reference number: A70809 MUID:98295987; pMID:9634230

A;Accession: A70869

A;Status: preliminary; nucleic acid sequence not shown; translation not shown A;Residues: 1-1660 <COL-A;Cross-references: UNIPROT:053215; UNIPARC:UPI00001652F8; GB:AL021246; GB:AL123456; NII A;Residues: strain H37RV
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                                     GAGGDGGDPGAG----GKGGAGGAGATEGVTGATGATVHSGGNGGKGGNGADATVAGANGG
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-SGAGAGSGVGVPGVGVPGVGVPG-----KGVPGVGP-----GVGP
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hypothetical glycine-rich protein Rv1450c - Mycobacterium tuberculosis (strain H37RV) C;Species: Mycobacterium tuberculosis C;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 20-Jun-2000 C;Accession: E70917 R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Gordon, Connoor, R.; Davles, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Holroyd, Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S. Nature 393, 537-544, 1998 R; Taylor, K.; Whitehead, S.; Barrell, B.G. A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G. A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete genor A;Reference number: A70500; MUID:98295987; PMID:9634230 A;Accession: E70917 A;Accession: E70917 A;Accession: E70917 A;Accession: E70917 A;Accession: Deciphering tubercalcaled sequence not shown; translation not shown A;Maccession: E70917 A;Accession: PMID:9634230 တ

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A;Molecule type: DNA
A;Residues: 1-1329 <COL>
A;Cross-references: UNIPARC:UPI00000C158B; GB:Z95844;
A;Experimental source: strain H37Rv
C;Genetics:
A;Gene: Rv1450c
C;Superfamily: collagen alpha 2(I) chain; fibrillar co GB:AL123456; NID:g3250713; PIDN: CF

Query Match Local al Similarity 33.8 340; Conservative 24.2%; Score 1010.5; DB 2; 33.8%; Pred. No. 4.4e-54; tive 35; Mismatches 382; collagen carboxyl-terminal Indels 249; Length 1329; Gaps 47 homology

181 GAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGS 240 134 291 177 41 GVGPGVGPGAGAGS-----GAGAGSGA-----GHGAEGVAGAAGEDLSPHGTSGGVGGDAGDGGTGGRGGWLAGAGGAGGAGGVGGTGGAGG G---GVGGTGGAGGLGGTLYGAGGHGGAGGPGPIGGVGGHGGVGGAAGLLGVGGHG-GAG GAGAAGGAGGSGGWLLGNGGVGGAGGQSLLGGATGGAGGNAGLFGVGGTGGPG-GPG-GP GAGAGSGAGAGSG--VGVPGVGVPG------GKGVPGVGPGVGPGVGP----GVGPGAGAG--AGFSRALIVAGDNGGDPGAGGAG-GTGGAGSTIGAHGAAGASPTSGGNGGAGGNGAHFSS VG-----VPG--VGVPGVGVPGKGVPGVGPGVGPGVGPGAGAGSGAGA-----GS -----VGVPGKGVPGVGPGVGP -GAGSGAGAGSGVGVPGVGVPGVGVP ----SGAGAGSGAGAGSGAGAGSG 133 350 180 290 40 87 234

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hypothetical glycine-rich protein Rv3507 - Mycobacterium tuberculosis (strain H37RV C;Species: Mycobacterium tuberculosis
C;Species: Mycobacterium tuberculosis
C;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 20-Jun-2000
C;Accession: E70806
R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Gord, Connor, R.; Davies, R.; Deviln, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Holroy Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S. Nature 393, 537-544, 1998
A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G. A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete ge A;Arcession: E70806
A;Status: preliminary; nucleic acid sequence not shown; translation not shown A;Arcession: E70806
A;Cross-references: UNIPARC:UPI0000D3AE7; GB:AL022022; GB:AL123456; NID:g3261554; A;Cross-references: UNIPARC:UPI0000D3AE7; GB:AL022022; GB:AL123456; NID:g3261554; A;Genetics:
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                                                     GNGGNG-GNGGDGGNGAHGTVGAQFVPATSLPTPNGGAGGNGGTGSNGGAPGPAGAPGPT
                                                                                  PGVGPGVGPGAGAGSGAGAGSGA------GAGSGAGAGSGVGVPG-VGVPGVG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GVGPGVGPG----
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                                                                                                                                                                       -GIGAQGTTITVPGNGGNAGDG-----
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RESULT 15
A70334
R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, I Mycobacterium tuberculosis (strain D.; Gordon, တ

695	GPGVGPG-VGPGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVG	, 639	ð
954	GGNGGNGGNGAAGGNGGNGGNGSLRGSGGAGGHGGNG-GNASRGMGGDGGTG	896	밁
638	GSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGV	, 599	ð
895	PGDGGNGANGSVFTNNGIGGNG-GNGGNAGPSGAGGSGGAGSTFGATGSSSSIHVN	84	문 5
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787	GHPGGTGGVGGTGGKGGDGGDGGAAPNGVAGSQGPGGAGGDGGTGGVGGNGGRGIDGADG	728	밁
499	AGABGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGA	445	S
727	NGSSAGA	66	문 :
444	VGPGVGPGVGPGVGPGAGAGSGAGAGSGAGAGSGAGAGSGV-GVPGVGVPGV	394	§
393	NGAPGNTPTSGGNGGRGADATGFGQTGASGGRGGDGGLVGNGSAGGAGGAGGAGGNGSKGLPG	608	문 5
607	GEGEAGGNG-GTFGNGGHGAAGALGVNGGVGGAGGHGGDFGVG-GAGGQGGSG	י ני	2 5
358	<	30	. 5
549	GDGGSGGAGSIGFNASAPGAAGSPGGNGGNGGPGGAGGEGGAGGLALAASGQNGSQG	493	D.
307	SGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGVGP	, 252	ð
492	QVGNGGAGGAGGAGGIGFGSEAPGRPGGAGGTG-GAGGNGGTQAGDG-GTGGAGGAG	435	뮍
251	GVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGAGAGAGAGAGAGAGAG	, 200	Ş
434		37	문 :
199	GVGPGVGPGVGPGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGV	15	ۇ
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315	GAGGAGGVLMGNGGNGGAGGAGGAGGAGGAGGAGGAGGNGGNGGNGGNGG	256	Д
96	GAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGP	, 51	ð
50	GAGA-GSGAGAGSGVGVPGVGVDGVGVDGVGVDGKGVPGVGPGVGPGVGPGVGPGVGPGAGAGAGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	, 1	음 성
44;	23.6%; Score 984; DB 2; Length 1306; Similarity 34.0%; Pred. No. 1.7e-52; 2; Conservative 34; Mismatches 386; Indels 224; Gaps	Query Match Best Local S Matches 332	
	1 source: strain H37Rv BC : collagen alpha 1(TV) chain	A; Experimental C; Genetics: A; Gene: Rv0578c C; Superfamily:	Ç,≯,Ç,≯
AL123456; NIC	:YPG: UNA 1-1306 <col/> :rences: UNIPROT:053775; UNIPARC:UPI00000D4FC8; GB:AL021942; GB:	Residues: 1-1306 Cross-references:	222
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; Holroyd, S.	.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares,	Connor, R jandream,	Z 72 '-

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Search completed: May 10, 2006, 10:44:30 Job time: 51 secs

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Q9VTR6_DROME
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Q8C9L8_MOUSE
ELN MOUSE
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Q5RKH4_RAT
ELN RAT
Q7VTB DROME
ELN BOVIN
PHXR5_MOUSE
C44358_NEPCL
Q8VIYY_MYCTU
Q8VIYY_MYCTU
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  06q294 agelenopsis
06py84 kukulcania
09biu8 argiope tri
017434 nephila cla
06fpr0 candida gla
09hw2 nephila ina
09hit2 nephila cla
09hw4 nephila cla
09wtr6 drosophila
09wtr6 drosophila
09bit7 nephila ina
09bit7 nephila ina
09u517 drosophila
094359 nephila cla
08c918 mus musculu
05rkh4 rattus norv
07yu48 dorosophila
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Result No.

4 5	44	43	42	41	40	39	9 U	37	36	35	34	33	32
992	998.5	1003	1006	1007	1007.5	1010.5	1017.5	1019	1034	1034	1041	1058.5	1064
23.8	23.9	24.0	24.1	24.1	24.1	24.2	24.4	24.4	24.8	24.8	24.9	25.4	25.5
1360	1274	1384	1381	1408	1408	1329	854	651	1665	1660	747	786	1468
N	N	N	N	N	N	N	N	N	N	N	Н	ب	N
Q7TWC4_MYCBO	Q6CHN8_YARLI	Q8VIZ1_MYCTU	Q6MWW9_MYCTU	Q7U022_MYCBO	Q8VK17 MYCTU	Q79FP2_MYCTU	Q9BIU4_DOLTE	Q9BIU9_ARGTR	Q7D721_MYCTU	Q79FD4_MYCTU	SPD1_NEPCL	ELN_HUMAN	Q9GUB5_GALME
Q7twc4 mycobacteri	Q6chn8 yarrowia li	Q8viz1 mycobacteri				Q79fp2 mycobacteri		Q9biu9 argiope tri			P19837 nephila cla		Q9gub5 galleria me

ALIGNMENTS

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PARTIAL NUCLEOTIDE SEQUENCE. MEDLINE-94365842; PubMed-716056; DOI=10.1007/BF00175878; MITA K., Ichimura S., James T.C.; "Highly repetitive structure and its organization of the silk fibroin gene."; J. Mol. Evol. 38:583-592(1994). [6] NUCLEOTIDE SEQUENCE OF 5179-5263, AND DISULFIDE BONDS. STRAIN-J-139;	Cell 16:425-436(1979). [4] PARTIAL NUCLEOTIDE SEQUENCE. STRAIN=Kinshu X Showa; MEDLINE=89094868; PubMed=3210244; MEDLINE=89094868; PubMed=3210244; Mita K., Ichimura S., Zama M., James T.C.; "Specific codon usage pattern and its implications on the secondary structure of silk fibroin mRNA."; J. Mol. Biol. 203:917-925(1988). [5]		NUCLEOTIDE SEQUENCE OF 1-168. NUCLEOTIDE SEQUENCE OF 1-168. MEDILINE-80045039; PubMed=498286; DOI=10.1016/0092-8674(79)90075-8; TBujimoto Y., Suzuki Y.; "The DNA sequence of Bombyx mori fibroin gene including the 5' flanking, mRNA coding, entire intervening and fibroin protein coding regions."; Cell 18:591-600(1979).	NCBL_TAXID=7001; [1] NUCLEOTIDE SEQUENCE. NUCLEOTIDE SEQUENCE: NEDIJINE=20330362; PubMed=10871375; DOI=10.1093/nar/28.12.2413; MEDIJINE=20330362; PubMed=10871375; DOI=10.1093/nar/28.12.2413; Zhou CZ., Confalonieri F., Medina N., Živanovic Y., Esnault C., Zhou CZ., Confalonieri F., Medina N., Perasso R., Li ZG.; Yang T., Jacquet M., Janin J., Duet M., Perasso R., Li ZG.; "Fine organization of Bombyx mori fibroin heavy chain gene."; Nucleic Acids Res. 28:2413-2419(2000).	

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                                                                                                                                                                         EMBL; AF226688; AAF76983.1; -; Genomic_DNA.
EMBL; V00094; CAA23432.1; -; Genomic_DNA.
EMBL; V00097; CAA23433.1; -; Genomic_DNA.
EMBL; S74439; AAB31861.1; -; mRNA.
EMBL; S74439; AAB31861.1; -; mRNA.
EMBL; X13869; CAA32076.1; -; mRNA.
EMBL; M35378; AAA27839.1; -; mRNA.
EMBL; AB017362; BAA33147.1; -; Genomic_DNA.
EMBL; CK538369; -; NOT_ANNOTATED_CDS; GENOMic_DNA.
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Takagi T., Mizuno S.;
"Determination of the site of disulfide linkage
light chains of silk fibroin produced by Bombyx
Blochim. Biophys. Acta 1432:92-103(1999).
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Tanaka K., Kajiyama N., Ishikura K., Waga S., Kikuchi A., Ohtomo K.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              - TISSUE SPECIFICITY: Produced exclusively in the posterior (PSG) section of silk glands, which are essentially modified salivary glands.

DOMAIN: Composed of antiparallel beta sheets. The strands of the beta sheets run parallel to the fiber axis. Long stretches of sill fibroin are composed of microcrystalline arrays of (-Gly-Ser-Gly-Ala-Gly-Ala-)n interrupted by regions containing bulkier residues. The fiber is composed of microcrystalline arrays alternating with amorphous regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FUNCTION: Core component of the same function: Core component of the same dhemically inert fiber.

and chemically inert fiber.

SUBUNIT: Silk fibroin elementary unit consists in a disulfident of silk fibroin elementary and a p25 glycoprotein in molar linked heavy and light chain and a p25 glycoprotein in molar linked heavy and light chain and a complex of approximately of 6:6:1. This results in a complex of approximately
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         European
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                                                                                                                                                            GLGAGL----
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   GAGSGSGAAA-
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Ampullate Silk cDNAs
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GSGAGLGRGAGTGAGAGSGAGSGAGAGSGS
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RESULT 3
Q6PY84 KUKHI
ID Q6PY84;
PAC Q6PY84;
DT 05-UUL-2004 (T
DE Major ampullat
OS KUKULCania hib
OC Araneomorphae;
OX NCBI TaxID=268
RN NULCEOTIDE SEQ
RN NULCEOTIDE SEQ
RN NULCEOTIDE SEQ
RN Tian M., Liu C
RT "Analysis of M
RT Spiders.";
RL Biomacromolecu
DR EMBL; AY571308
FT NON TER
SQ SEQÜENCE 760
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Best Local
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last amontation update)
Major ampullate spidroin 1 (Fragment).
Kukulcania hibernalis (Southern house spider).
Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida;
Eukaryota; Metazoa; Arthropoda; Chelicerata; Kukulcania.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; AY571308; AAT08433.1; -;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Biomacromolecules 0:0-0(2004).
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"Analysis of Major Ampullate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE.
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 AGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVFGKGVPGVGPGVGPGVGPGVGPGAG
                               RRGPGGARSRGGAGAGAGAGSGVGGYGSG-SGAGA-----
                                                                                             AG----
                                                                                                                           PGVGVPGVGVPGKG------VPGVGPGVGPG-VGPGVGPGAGAGSGAGAGSG
                                                                                                                                                                                                                     PGVGVPGVGVPGXGVPGVGPGVGPGVGPGAGAGSGAGAGSGAGAGSGAGAGSGVGV
                                                                                                                                                                                                                                                                                      AVVFESAGPGEEAGSSGDGASAAASAAAAAGAGSGRRGPGGARSRGGAGAGAGAGSGVGG
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                                                                                                                                                                                        PGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGSGAGAGSGAGAGSGVGV
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760 AA;
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ilarity 46.3%;
Conservative 3
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                                                                                            GRGRGGRGGEAFSASSASSAVVFESAGPGEEAGSSGGGASAAASAAAAAGAGSG
                                                            ----AGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAG
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                                                                                                                                                        -GSGAGAGAGAGFGVGAGAGAGAGFGSGAGAGSGAGAGYGAGR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 1531; DB 2; Length 760;
Pred. No. 1.5e-84;
37; Mismatches 224; Indels 204;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Silk cDNAs from Two Non-Orb-Weaving
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                             -GAGAGAGAGGEGGFGEGQG
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RESULT 4
Q9B1U8 ARGTR PRELIMINARY;
AC Q9B1U8;
AC Q9B1U8-2003 (TrEMBLrel. 17,
DT 01-JUN-2003 (TrEMBLrel. 12,
DE Flagelliform silk protein (1,
DE Name=Plag;
AC NCBL_TAXID=156845;
AN NUCLEOTIDE SEQUENCE.
AC MEDLINE=21179804; PubMed=11:
AC GACESY J., HAYAShi C., Motr.
RT FIDIOIN SEQUENCE.
RY MEDLINE=21179804; Conserv.
RT Fibroin sequences.";
RL Science 291:2603-2605 (2001)
RN NUCLEOTIDE SEQUENCE.
RA GACESY J.E., HAYASHI C.Y.;
RL Science 291:2603-2605 (2001)
RN NUCLEOTIDE SEQUENCE.
RA GACESY J.E., HAYASHI C.Y.;
RL Science 291:2603-2605 (2001)
RN NUCLEOTIDE SEQUENCE.
RA GACESY J.E., HAYASHI C.Y.;
RL Submitted (FEB-2001) to the
DR EMBL; AF350265; AAX30594.1;
DR GO; GO:0005817; P:phosphate
DR GO; GO:0006817; P:phosphate
DR INCEPTO; IPRO08160; Collagy
FT NON_TER 1002
AA; 75889 1
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Best Local S
Matches 392
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01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Flagelliform silk protein (Fragment).
Name=Flag;
                                                                                                                                                                                                                                                                                                                                                                                       Gatesy J.E., Hayashi C.Y.;

Submitted (FEB-2001) to the EMBL/GenBank/DDBJ

EMBL; AF350265; AAX30594.1; -; mRNA.

GO; GO:0005737; C:cytoplasm; IEA.

GO; GO:0006817; P:phosphate transport; IEA.

InterPro; IPR008160; Collagen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=21179804; PubMed=11283372; Gatesy J., Hayashi C., Motriuk D., "Extreme diversity, conservation, a fibroin sequences."; Science 291:2603-2605(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Argiope trifasciata (Banded garden spider).
Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Ara
Araneomorphae; Entelegynae; Araneoidea; Araneidae; Argiope.
NCBI_TaxID=156845;
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                                                                                                                                                                                                                                   Mismatches
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01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Minor ampullate silk protein MiSpl (Fragment).
Mephila clavipes (Orb spider).
Eukaryota, Metazoa, Arthropoda; Chelicerata; Arachmida;
Araneomorphae; Entelegynae; Araneoidea; Tetragnathidae;
                                          Colgin M.A., Lewis R.V.;
"Spider minor ampullate silk proteins contain new and highly conserved non-silk-like 'spacer regions protein Sci. 7:667-672(1998).
  EMBL; AF027735;
NON_TER 1
                                                                                                                                                                                                       NCBI_TaxID=6915;
                                                                                                                                          MEDLINE=98200471; PubMed=9541398;
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                                                                                                                                              PGVGVPGVGVPGVGVGVGPGVGPGVGPGAGAGSGAGAG
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Al Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla B.,
Al Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V.,
Barnay S., Blanchin S., Beckerich J.-M., Beyne E., Bleykasten C.,
Al Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,
Al Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
Al Hantraye F., Hennequin C., Jauniaux N., Joyet P., Kachouri R.,
Al Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,
Al Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
Al Pellenz S., Potler S., Richard G.-F., Straub M.-L., Suleau A.,
Al Pellenz S., Potler S., Richard G.-F., Straub M.-L., Suleau A.,
Swennen D., Tekaia F., Weeolowski-Louvel M., Westhof E., Wirth B.,
Zeniou-Meyer M., Zivanovic Y., Bolotin-Fukuhara M., Thierry A.,
Bouchier C., Caddron B., Scarpelli C., Gaillardin C., Weissenbach J.,
Mincker P., Souciet J.-L.,
Mincker P., Straub M., Mincker P., Souciet J.-L.,
Mincker P., Souciet J.-L.,
Mincker P., Souciet J.-L.,
Mincker P., Souciet J.-L.,
Mincker P., Straub M., Mincker P., Souciet J.-L.,
Mincker P., Souciet J.-L.,
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Similarities with tr|Q08294 Saccharomyces cerevisiae
OrderedLocusNames=CAGLOJ01771g;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Candida glabrata (Yeast) (Torulopsis glabrata).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; mitosporic Saccharomycetales; Candida.
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PubMed=15229592; DOI=10.1038/nature02579;
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AGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSG
                                                                                              GAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGS
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Pred. No. 2.6e-76;
2; Mismatches 215;
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RESULT 7
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Matches 412
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Q9NHW2;
Q9NHW2;
Q1-OCT-2000 (TrEMBLrel. 15,
01-OCT-2000 (TrEMBLrel. 24,
01-JUN-2003 (TrEMBLrel. 24,
Flagelliform silk protein ()
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene.";

Science 287:1477-1479(2000).

EMBL; AF218633; AAF36091.1; -; Genomic_!
GO; GO:0005737; C:cytoplasm; IEA.
GO; GO:0006817; P:phosphate transport;
InterPro; IPR008160; Collagen.

1884
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Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Araneae;
Araneomorphae; Entelegynae; Araneoidea; Tetragnathidae; Nephila.
NCBI_TaxID=115969;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE.
MEDLINE=20156766; PubMed=10688794;
Hayashi C.Y., Lewis R.V.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Molecular architecture and
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       VGP-GVGP-GVGPG-----
                                                                                                  GPGGSGPGGAGSGGVGPGGYGPGGAGPG-GAGPGGAGPGGAGPGGAGTGGAGTGG
                                                                                                                                      GPG----AGAGSG------AGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPG
                                                                                                                                                                                                     PGGAGPGGYGPGGAGPG-
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1884 AA;
                                                                                                                                                                                                                                                                                                                                                                                           32.7%; Score 1365; Dilarity 42.3%; Pred. No. 2.6e Conservative 16; Mismatches
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(TrEMBLrel. 15, Last sequence update)
(TrEMBLrel. 24, Last annotation update)
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RESULT 8

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DT 01-AUG-1988 (
DT 10-MAY-2005 (
DE Elastin precu
GN Name-ELN;
OS Gallus gallus
OC Eukaryota; Me
OC Archosauria;
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NUCLEOTIDE SEQUENCE.
MEDLINE=87242320; PubMed=3593675;
Bressan G.M., Argos P., Stanley K.K.;
"Repeating structure of chick tropoelastin
                                                                                                                                                                                                                                                                    01-AUG-1988 (Rel. 08, Created)
01-AUG-1988 (Rel. 08, Last sequence
10-AMY-2005 (Rel. 47, Last annotatic
Elastin precursor (Tropoelastin) (Fr
                                                                                                                                                                                                Gallus gallus (Chicken).
Eukaryota; Metazoa; Chor
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EMBL; M21880; AAA49082.1; -;
EMBL; M15889; AAA49108.1; -;
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Baule V.J., Foster J.A.;
"Multiple chick tropoelastin mRNAs.";
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Biochemistry 26:1497-1503(1987).
[2]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FUNCTION: Major structural protein of tissues such as aorta a nuchal ligament, which must expand rapidly and recover comple SUBUNIT: The polymeric elastin chains are cross-linked togeth into an extensible 3D network.

SUBCELIULAR LOCATION: Extracellular matrix of elastic fibers.

ALTERNATIVE PRODUCTS:

Event=Alternative splicing; Named isoforms=2;

Comment=Additional isoforms seem to exist;
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RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N., RA Brandon R.C., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., RA Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D., RA Man K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G., RA Abril J.F., Agbayani A., An H.-J., Andrews Pfannkoch C., Baldwin D., RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P., Rockies K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I., Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P., RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P., RA Goldek A., Gong F., Gorrell J.H., Gu Z., Gelbart W.M., Glasser K., Andrew B.C., Gabriellan A.E., Garg N.S., Gelbart W.M., Glasser K., Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M., Alakin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A., RA Kimmel B.E., Kodira C.D., Kraft C., Kravizz S., Kulp D., Lai Z., Lang Y., Lin X., RA Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.L., RA Lasko P., Levitsky A.A., Li J.H., Li Z., Liang Y., Lin X., RA Merkulov G., Milshina N.V. Mobarry C., Morris J., Moshrefi A., RA Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M., RA Palazzolo M., Pittman G.S., Pan S., Pollard J., Pacleb J.M., RA Ra Rainert K., Remington K., Saunders R.D.C., Scheeler F., Shen H., RA Ra Spier E., Sten H., Sanders R., Dector C., Turner R., Venter E., Wang A.H., Wang X., RA Mang Z.-Y. Wassarman D.A., Weinstock G.M., Weissenbach J., Ra Mang S., Yao Q.A., Shith H.O., Shith H.O., Smith H.O., Smit
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01-OCT-2002 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
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Amanatides P.G.,
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Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CG5700-PB.
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Ephydroidea; Drosophilidae; Drosophila.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PubMed=10731132; DOI=10.1126/science.287.5461.2185;
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Best Local
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FlyBase; FBgnuvaura, pro.
FlyBase; FBgnu028573; pro.
GO; GO:0005578; C:extracellular matrix (sensu Meta GO; GO:0007507; P:heart development; IMP.
GO; GO:0002009; P:morphogenesis of an epithelium; InterPro; IPR009765; Pericardin rpt.
Pfam; PF07054; Pericardin rpt; 34.
Pfam; PF07054; Pericardin rpt; 34.
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Science 287:2185-2195(2000).
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Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S.,
Hradecky P., Huang Y., Kaminker J.S., Millburn G.H., Prochnik S.E.,
Smith C.D., Tupy J.L., Whitfield E.J., Bayraktaroglu L., Berman B.I
Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Berkeley Drosophila Genome Project;
Celniker S., Carlson J., Wan K., Pfeiffer
Hoskins R., Stapleton M., Pacleb J., Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=22426070; PubMed=
Kaminker J.S., Bergman C.
                                                                                                                                                                                                                                                                                                                                                Ensembl;
                                                                                                                                                                                                                                                                                                                                                                 EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           "Drosophila melanogaster
Submitted (MAR-2000) to t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "The transposable elements of the Drosophila melanogaster genomics perspective.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lewis S.E.;
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                                                                                                                                                                                                                                                                                                                             AE003543; AAF49980.2; -; Genomic_DNA.bl; CG5700; Drosophila melanogaster.se; FBgn0028573; CG5700.
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                                                                                                                                                          415;
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                                                                                                                                                                         Similarity
                              GQTGAGQPGYGSQPVVGAQTGTGQPGYGAQPGVGTQTGAGQPGYGSQPGIGGQTGARQPG
                                                                                            GYGSQLGVGAQAGAGQPGYGDQPGVGTQPGVGQPGYGSQPGTGGQTGAGQPGYGSQPGTG
 ----PGVG--PGVG-PGAGAGSG----AGAGS-GAGAGSGAGAGSGVGVPGVGV-PGVGV
                                                             SGAGAGS-GAGAGSGAGAGSGVGVPGVGV-PGVGV-PGKGVPGVG--PGVG---
                                                                                                                        GAGAGSGAGAGSGVGVPGVG-VPGVGV-PGKGVPGVG--PGVGPGVG---PGVGPGAGAG
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                                                                                                                                                          Conservative
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Celniker S.
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PubMed=12537573;

PubMed=12537573;

Lewis S.E.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                             lease 4 sequence.";
EMBL/GenBank/DDBJ
                                                                                                                                                                       Score 1312.5; DB 2; Pred. No. 3.4e-71;
                                                                                                                                                            Mismatches
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Svirskas F
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Smith E.,
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RESULT 10
Q9BIT2 PLETR
ID Q9BIT2;
ID Q9BIT2;
AC Q9BIT2;
DT 01-JUN-2001 (TREMBLrel. 17, Created)
DT 01-JUN-2001 (TREMBLED 17, Last sequence update)
DT 01-UN-2001 (TREMBLED 17, Last sequence update)
DT 01-OTCT-2002 (TREMBLED 122, Last annotation update)
DE Fibroin 1 (Fragment).
OS Plectreurys tristis (Spider).
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MEDLINE=21179804; PubMed=11283372;
Gatesy J., Hayashi C., Motriuk D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Araneae;
Araneomorphae; Haplogynae; Plectreuridae; Plectreurys.
NCBI_TaxID=33319;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               fibroin sequences.";
Science 291:2603-2605(2001).
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   GAGGAGAGFGSGLGLGYGVGLSSAQAQAQSAAAARAQADAQAQAYAAAQAQAQAQAQAQAQA
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                                                                                                                                             -AGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVPGK
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DR NUCLEOTIDE SEG
RX MEDLINE=201567
RA MAYASHI C.Y.,
RT "MOLECULAR ARCHAEL
RY MEDLINE=201567
RA HAYASHI C.Y.,
RT "MOLECULAR ARCHAEL
DR GO; GO:0006817
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Science 287:1477-1479(2000).
Science 287:1477-1479(2000).
EMBL; AF218621; AAF36090.1; -; Genomic_DNA.
GO; GO:0005737; C:cytoplasm; IEA.
R GO; GO:0006817; P:phosphate transport; IEA.
InterPro; IPR000160; Collagen.
InterPro; IPR000209; Pept S8_S53.
PR INTERPRO; IPR000209; Pept S8_S53.
PR PROSITE; PS00138; SUBTILASE_SER; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 418; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nephila člavipes (Orb spider).
Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Araneae;
Araneomorphae; Entelegynae; Araneoidea; Tetragnathidae; Nephila.
NCBI_TaxID=6915;
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01-OCT-2000 (TYEMBLrel. 26, Last sequence update)
01-WAR-2004 (TYEMBLrel. 26, Last annotation update)
Flagelliform silk protein (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=20156766; PubMed=10688794;
Hayashi C.Y., Lewis R.V.;
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 GP-GVGPGAGAGSGAGAGS----GAGAGSGAGAG----SGVGVPGVGVPGVGVPGKGVP-
                               GGAGGAGGSGGTTIVEDLDITIDGADGPITISEELTIGGAGAGGSGPGGAGPGNVGPGRS
                                                                                          PGGAGTSGAGPGGAGPGGAGPGGAGRGGAGRGGVGRGGAGRGGARGAGGA
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2249 AA;
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Pred. No. 1.6e-70;
B; Mismatches 342;
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Submitted (FEB-2001) to the EMBL
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Best Local Similarity
Matches 406; Conserv
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EMBL; AF203342; AAF11280.1; -; mRNA.

FlyBase; FBgn0028573; pcc.

GO; GO:0005578; C:extracellular matrix (sensu Me: GO; GO:0007507; P:heart development; IMP.

InterPro; IPR009765; Pericardin rpt.

Pfam; PF07054; Pericardin rpt; 33.

SEQUENCE 1729 AA; 164661 MW; DA9B1B7FABSCDED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN=Oregon R; MEDLINE=22065272; PubMed=12070098; MEDLINE=22065272; PubMed=12070098; Chartier A., Zaffran S., Astier M., Semeriva M., Gratecos D.; "Pericardin, a Drosophila type IV collagen-like protein is involute morphogenesis and maintenance of the heart epithelium during dorsal ectoderm closuce.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Name=prc; ORFNames=CG5700;
Drosophila melanogaster (Fruit fly).
Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Miscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAY-2000 (TrEMBLrel. 13, 01-MAY-2000 (TrEMBLrel. 13, 01-MAR-2004 (TrEMBLrel. 26,
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(7_DROME PRELIMINARY;
Q9U617_DROME PRELIMINARY;
Q9U617;
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-----PGVG---PGVG-------PGAGAGSGAGAGSGAGA---GS--GAGAGSGVGVPGVG
                                                                           GSQPGIGGQTGAGQPGYGTQPGIGGQTGAGQPGYGTQPGIGAQTGAGQPGYGTQPGIGGQ
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CO Nephila
OS Nephila
OS Nephila
OC Eukaryott
OC Araneomo
OX NCBI_Tax
RN (1]
RP NUCLEOTI
RX MEDLINE-
RA Hayashi
RT "Evideno
RT elastici
RT "Buideno
RT elastici
RT EMBL, AF
                                                                                                                                                                                                                        O44359_NEPCL PRELIMINARY; PRT; 907 AA.

C 044359;
C 044359;
C 01-JUN-1998 (TrEMBLrel. 06, Created)
C 01-JUN-1998 (TrEMBLrel. 06, Last sequence update)
C 01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Flagelliform silk protein (Fragment).

Namme=Flag;
Nephila clavipes (Orb spider).
C Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Ara Araneomorphae; Entelegynae; Araneoidea; Tetragnathidae; Nephila Clavipes (Orb spider).
NUCLEOTIDE SEQUENCE.

MEDLINE-98153262; Pubmed=9480768; DOI=10.1006/jmbi.1997.1478;
Hayashi C.V., Lewis R.V.;
"Evidence from flagelliform silk cDNA for the structural basi
"Estidence from flagelliform silk cDNA for the structural basi
elasticity and modular nature of spider silks.";
J. Mol. Biol. 275:773-784(1998)
EMBL; AF027973; AAC38847.1; -; mRNA.
GO; GO:0005737; C:cytoplasm; IEA.
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                        GAGAGSGAGAG
                                                                      SGAGAGSGAGAGSGVGVPGVGVPGVGVPGVGVPGVGP-GVGPG-----VGPG-----VGP
                                                                                                DLDITIDGADGPITISEELTISGAGGSGPGGAGPGGVGPGGSGPGGVGPGVGPG
                                                                                                                                                GGYGPGGSGAG-GTGPGGAGGAG-GAGGSGGAGGSGGAGGSGGAGGSGGVGGSGGTTITE
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 791
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RC STRAINS—22354683; PubMed=12466851; DOI=10.1038/nature01266;

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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
10-MAY-2005 (TrEMBLrel. 30, Last annotation update)
10-MAY-2005 (TrEMBLrel. 30, Last annotation update)
Mus musculus 3 days neonate thymus cDNA, RIKEN full-length enriched
library, clone:A630042I19 product:elastin, full insert sequence
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Tennam M. Schmences "...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=FVB/N; TISS
MEDLINE=22388257;
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"Analysis of the mouse transcriptome based 60,770 full-length cDNAs.";
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PubMed=12477932; DOI=10.1073/pnas.242603899;
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Strausberg R.; Submitted (APR-2003) to the EMBL; AK041860; BAC31084.1; EMBL; BC051649; AAH51649.1;

EMBL/GenBank/DDBJ databases

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GO; GO:0030833; P:regulation of actin filament polymerization; IMP.
GO; GO:0043149; P:stress fiber formation; IMP.
InterPro; IPR003979; troppelastin.
PRINTS; PR01500; TROPOELASTIN.
PRINTS; PR01500; TROPOELASTIN.
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SELPAK po
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Silk-elas
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Silk/elas
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76.3	76.5	77.9	78.0	78.0	78.2	78.2	78.2	78.7	78.7	79.0	79.0	79.0	79.0	79.0	79.0	79.0	79.2	79.2	79.2	79.2
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Aar80253	Abp53481	Adu81676	Abp53473	Aar80252	Aeb57388	Adu81677	Abp53466	Aaw49724	Aar80335	Ade44977	Abg69272	Aay78282	Aaw53523	Aaw26347	Aar41012	Aap82961	Aeb57386	Adu81671	Adu98740	Adr70478
Polymer S	Protein p	Cotton bi	Protein p	Polymer S	SELP47K-P	Radioprot	SELP8 rel	Protein p	Protein p	Recombina	Silk/Elas	SELP3 ami	Amino aci	SELP3 syn	SELP3 mul	SEPL3 pro	SELP 67K	Silk/elas	Active ag	Silk-elas

ALIGNMENTS

RESULT 1 ADX51951 ADX51951 ADX51951 ADX ADX5 AX ADX5 AX ADX5 AX repe XX repe XX repe XX repe XX W02C XX 04-L XX 04-L XX (DO) PA (GEN XX (DO) PA (GEN XX WPI) PT FORT PT FORT PT Spet XX Clai XX Clai XX Clai XX Clai XX Seq 04-DEC-2003 Repeat 06-MAY-2004 ADK51951 standard; protein; 780 AA 20-MAY-2002; 2002US-0381913P. Unidentified ADK51951; 20-MAY-2003; 2003WO-US015757. WO2003099465-A1 repeat protein. (DOWO) DOW CORNING (GEMV) GENENCOR INT protein polymer repeat sequence, DOW CORNING CORP. (first entry) SEQ Ħ

19

Forming inorganic structure, for use as e.g. electronic composite materials, comprises providing substrate with polymer, and exposing substrate to precursor containing Claim 12; Page 8; 27pp; species. English. c photonic and na h repeat protein g inorganic

WPI; 2004-023263/02.

Kumar M;

The present invention relates to a method for forming an inorganic structure. The method comprises providing substrate with a repeat protein polymer, and exposing the substrate to a precursor comprising inorganic species. The repeat protein polymer catalyzes the formation of an inorganic structure on the substrate. The repeat portion of the repeat protein polymer may be ADKS1951-ADKS1957, which may have a head sequence at the N-terminal end (ADKS1958) and a tail sequence at the C-terminal

Sequence 780 AA;

end. (ADK51959-ADK51966).

Length

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RESULT 2
AD70468
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Matches 780; Conserv
personal care composition; repeat sequence protein polymer; hair care composition; shampoo; conditioner; anti-dandruff treatment; styling aid; styling conditioner; hair repair treatment; chemical hair treatment; hair dye; skin care composition; body wash; antimicrobial cleanser; skin protectant cream; body lotion; facial cream; sturising cream; facial cleansing emulsion;
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anti-acne treatment; facial toner; exfoliating cream; facial mask; after shave balm; sunscreen; anti-fungal treatment; skin protectant; antiperspirant; cosmetic composition; makeup; eye gel; high-melting point lipstick; lipstick; lip gloss; lip balm; mascara; eyeliner; pressed powder formulation; foundation; nail care composition; nail enamel; cuticle treatment; nail polish; nail treatment; polish remover; oral care composition; toothpaste; mouth rinse; breath freshener; teeth whitening treatment; over-the-counter pharmaceutical composition; transparent film formation; hydrogel formation; water-solubility; luster; softness; moisture retainment; tensile property; viscoelastic behaviour; glass translion temperature; cloud temperature; glass translion temperature; silk; elastin; repeating sequence unit.
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12-MAR-2004; 2004US-00800179

12-MAR-2003; 2003US-0454077P.

(KUMA/) KUMAR N ε

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Kumar M,

WPI; 2004-675584/66

Personal care composition useful as cosmetic, hair care or skin care product, comprises repeat sequence protein polymer and compounds such carriers, excipients, liposomes, active ingredients, or emollients. 86

Example 1; SEQ ID NO 19; 50pp; English.

The invention describes a personal care composition (I) comprising a composition protein polymer with the balance of the composition composition comprising and one or more compounds chosen from carriers, excipients, liposomes, active ingredients, biological or botanical products, conditioner, ancidents, thickening agents, silicone components, organic sunscreens, preservatives, neutralizing agents, competitioner, anti-dandruff treatment, styling aids, styling accompositioner, hair repair or treatment, serum, lotion, cream, ponded, or chemical treatment such as hair dye; skin care composition such as conditioner, hair repair or treatment, serum, lotion, cream, facial protectant cream, body vash, body wash, antimicrobial cleanser, skin collating gent, skin core composition such as collaransing emulsion, surfactant-based facial cleanser, facial exfoliating gel, anti-acne treatment, facial toner, exfoliating cream, facial mask, core-the-counter drugs comprising anti-fungal treatments, anti-acne composition chosen from eye gel, high-melting point comprising a makeup composition chosen from eye gel, high-melting point composition, lip gloss, lip balm, mascara, eyeliner, pressed composition. The composition is a mascara, eyeliner, pressed composition. The composition is a mascara such as nail care composition such as nail composition is a mascara such as non-waterproof mascara, waterproof mascara, or valumising mascara, water based mascara, or ceyelash or eyebrow treatment; and over-the-counter pharmaceutical composition water in-silicone foundation, or such as aloose composition such as a shail care composition such as a shail composition is a mascara water-based mascara, or ceyelash or eyebrow treatment; a pressed powder foundation such as sacing curling mascara, anhydrous waterproof mascara, water-based mascara, or ceyelash or eyebrow treatment; a pressed powder foundation such as such as aloose composition such as transition, but a such as a core composition such as transition, ce composition such as transition, an

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Best Local :
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 Active agent
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention relates to a novel system for providing controlled release delivery of an active agent which comprises a repeat sequence protein polymer and at least one active agent. The invention may be useful in a hair care composition, skin care composition, nail care composition, cosmetic composition, oral care composition or over-the-counter pharmaceutical composition. The system may be useful in shampos, gels, mousses and other hair care products, rinse-off conditioners, skin care products such as moisturizers, toners and makeup and nail care products such as polishes and polisher smovers. The invention enables controlled release of active agents. The present sequence is that of an SELP47K protein which may be used in the system of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           System for providing controlled release delivery of active agent useful for incorporating active agents into personal care product compositions, comprises repeat sequence protein polymer and active agent.
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Chimeric.
The invention relates to biomolecular conjugates comprising a repeat sequence protein polymer (i.e., an engineered protein) and at least one active agent, such as silicones, fragrances, dyes, tints, UV actives, sunscreens, lanolin, vitamins, bleaches, thickening agents, antimicrobials and biologically active polypeptides. The repeat sequence protein polymer preferably comprises a repeating amino acid sequence unit derived from elastin, collagen, abductin, byssus, flagelliform silk, dragline silk, gluten high molecular weight subunit, titin, fibronectin,
                                                                                                                                                                                                                                                                                                               Biomolecular conjugates useful in personal care products e.g. hair composition, commetic, oral care composition, comprising conjugation product of repeat sequence protein polymer and active agents.
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                                                                                                                                                                                                                                                                Claim
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CC arepeat sequence protein polymer (RSPP) or its fragment, and a carrier correction, where repeat sequence protein polymer provides at least one commercic benefit; or (b) a RSPP or its fragment, and one or more commercic benefit; or (b) a RSPP or its fragment, and one or more commonds chosen from carriers, excipients, liposomes, biological or botanical products, and where repeat sequence protein polymer or its commental damage, where the environmental damage comprises damage due to exposure to radiation (especially ultraviolet light) or damage due to exposure to chemicals or free radicals. (I) is useful as a skin care composition such as moisturizing body washes, body washes, antimicrobial composition skin protective creams, body lotions, facial creams, composition composition facial masks, aftershave balms, composition composition facial masks, aftershave balms, composition facial treatments, skin protectants, anti-perspirants, or radioprotective such as sunscreen chosen from non-water-resistant sunscreens. (I) is useful as a sunscreens, and water-in silicone sunscreens. (I) is useful as a composition such as eye gels, anti-string protective such case sunscreens. (I) is useful as a sunscreens and water-in silicone sunscreens. (I) is useful as a sunscreens chosen from loose powders, blushes, eye shadows, and communications chosen from loose powders, blushes, eye shadows, and water-in-oil foundations, water-in-oil foundation
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Best Local S
Matches 752
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Protein polymers comprising repeating units and enzyme-catalysed covalent bond formation useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pendent group; repeating unit; enzyme recognition site; sealant; fibrin; enzymatic cross-linking; biocompatible material; structural integrity; medical adhesive; wound closure; tissue repair; transglutaminase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 7; Page 65; 138pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   material for wound closure and tissue repair.
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Pred. No. 6.9e-217;
0; Mismatches 26;
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                                                                                                                                       Misc-difference
                                                                                                                                                                               Peptide
                                                                                                                                                                                                                    Region
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                                                                                                                                                                                                                                                                                                   Region
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                                                                                                                                                                                                                                                                                                                                                                   This SELPBK polymer sequence (mol.wt. 69,772) has been constructed from monomers composed of repeat units derived from silk fibroin (AAW09206) and elastin (AAW09207) (claim 4). The monomer (AAW09212) has been amplified by multicopy cloning in Escherichia coli (using plasmid pPT0345) to produce a recombinant protein polymer backbone with Lys functional groups for subsequent polyfunctional chemical crosslinking. The crosslinked polymer product forms a strongly adherent tissue adhesive or sealant. The polymer may be used to seal defects in vessel walls, e.g. artery, vein, capillary, lung, dura or colon, to increase tissue mass, or to produce biocompatible films for in vivo use. The sealants have the biocompatibility of fibrin glues, but set more quickly, and have greater strength. They are readily prepared by recombinant methods, are easy to administer, and are gradually resorbed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tissue repair adhesive comprising polymer of structural protein repeat units - contg. hetero-atom functional gps. reactive with crosslinking agent, combines biocompatibility and high bonding strength.
                                                                                                                                                                                                                                                                                                                                                 Sequence 884
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GAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVG-PGVG-PGVG-PGA 170
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Pred. No. 6.9e-217;
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RESULT 8
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                                                                                                                                                                                                            Preparation of synthetic repetitive DNA - useful for construction large protein polymers having repeating units, used in structural material, e.g. synthetic silk.
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                                                                                                                                                                      Example 7; Page 89; 127pp;
                                                                                                                                                                                                                                                                                                                                               Ferrari
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This is the amino acid sequence of pPT0345 comprising the SELPBK protein, used in the method of invention, which involves the preparation of synthetic DNA sequence having repeating units from about 3-15 codons and encoding a protein of at least about 30 kDa. The method is useful for the production of high molecular weight polymers (e.g. synthetic silk), either nucleic acids or peptides that are the expression products of the

and particularly high molecular weight peptides

containing

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RESULT 9
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Pred. No. 6.9e-217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SELPBK polymer is a synthetic silk-like protein comprising multiple CC copies of a monomer (see AAW49726) consisting of repeating units of silk-clike sequences (GAGAS) and elastin-like sequences (VPGVG) including a CC reactive lysine residue. It was expressed in Escherichia coli from CC plasmid ppT0345 as a 80 kDa protein. The SELPBK polymer was specifically CC designed to be functionalised with different reagents reactive to amines. CC claimed recombinant protein polymers are capable of covalent crosslinking by enzymatic reaction to form products which set quickly and have good adhesive properties and high strength. The proteins can comprise a CC repetitive amino acid backbone of repetitive units having collagen, CC fibroin, elastin or keratin motifs and at least 2 enzyme recognition consequences comprising a glutamine or lysine capable of enzyme catalysed CC sequences comprising a glutamine or lysine capable of enzyme catalysed CC sequences comprising a glutamine or lysine capable of enzyme catalysed CC sepetide formation. The products can be used as medical adhesives and CC sequences in the closure of wounds and repair of damaged tissue. (Updated CC on 25-MAR-2003 to correct PF field.)
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Matches 752
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                                                                                      Conservative
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healing; transglutaminase; cross-linking.
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Pred. No. 6.9e-217;
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This invention describes a novel crosslinked protein composition (A) which, before crosslinking, the protein (I) is new and comprises at I 70 wt.% of repeating units GAGAGS (I) and GVGVP (2), and in at least repeating units an amino acid (aa) is substituted by Lys or Arg to provide a Lys/Arg equivalent weight of 1-20 kD. (I) contains at least
                                                                                                                                                                               Crosslinked protein composition, useful as tissue adhesive or comprises peptide repeating units that contain functional growwith crosslinker.
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RESULT 11 ABG31412 ID ABG31

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90.6%;
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Matches 752
                                                                                                                                                                                                                                                                                                                                  The present invention relates to protein polymers having repetitive units from naturally occurring structural proteins such as fibroin, elastin, collagen and keratin. The polymers comprise a functional group which can be chemically crosslinked with appropriate crosslinkers. The protein polymer is produced by recombinant DNA technology. The protein polymer is useful for sealing or filling a defect in viable tissue, particularly for augmenting tissue mass. The protein polymer is useful in a variety of applications related to their physical, chemical and biological properties, and/or to bond together separated tissue to provide a stable, flexible or resorbable bond. The protein is particularly useful as a sealant or adhesive, in wound healing or as a burn dressing e.g. to stop or staunch the flow of fluid (e.g. blood) through ruptured vessels (e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Crosslinked protein composition used as sealant or adhesive for sealing or filling defect in viable tissue, as burn dressing, or in wound healing e.g. to staunch flow of fluid e.g. blood, through ruptured vessels.
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02-MAY-1996;
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keratin; SELP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SELP8K polymer encoded by plasmid pPT0345
                                                                                                                                                                                                                                                                                            Sequence 884
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                                                                                                                                                                                           PROTEIN
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96US-00642246.
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Pred. No. 6.9e-217;
0; Mismatches 26;
                                                                                                                                                                                                                                                              Length 884;
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Crosslinked protein composition useful as sealing a defect in tissue,

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RESULT 12
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XX 05-AP
PF 05-AP
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02-MAY-1996;
29-NOV-1999;
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                                                                                                                                                                                                                                                                                                                          US2003104589-A1
                                                                                                                                                                                                                                                                                                                                                                 Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                       coating; vulnerary.
                                                                                                                                                                                                                                                                                                                                                                                                                            Crosslinked protein; sealant; depot; drug; therapy; gel; film; thread;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Plasmid pPTO345 SELP8K polymer protein.
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99US-00451206.
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Best Local
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                                                                                                                                                                                                                    VPGVG-PGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGV
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90.6%;
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Pred. No. 6.9e-217;
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RESULT 13
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KW hair care composition; shampoo; conditioner; anti-dandruff treatment;
KW styling aid; styling conditioner; hair repair treatment;
KW chemical hair treatment; hair dye; skin care composition; body wash;
KW antimicrobial cleanser; skin protectant cream; body lotion; facial cream;
KW moisturising cream; facial cleansing emulsion;
KW surfactant-based facial cleanser; facial exfoliating gel;
KW anti-acne treatment; facial toner; exfoliating cream; facial mask;
KW after shave balm; sunscreen; anti-tungal treatment; skin protectant;
KW anti-perspirant; cosmetic composition; makeup; eye gel;
KW wish-melring point lipstick; lipstick; lip gloss; lip balm; mascara;
KW eyeliner; pressed powder formulation; foundation; nail care composition;
KW moistn remover; oral care composition; toothpaste; mouth rinse;
KW breath freshener; teeth whitening treatment;
KW worer-the-counter pharmaceutical composition; transparent film formation;
KW worsture retainment; tensile property; viscoelastic behaviour;
KW stansition temperature; cloud temperature; softness;
KW decommosition temperature; cloud temperature; softness;
KW decommosition temperature; cloud temperature;
The invention describes a personal care composition (I) comprising a CC repeat sequence protein polymer with the balance of the composition CC comprising one or more compounds chosen from carriers, excipients, CC liposomes, active ingredients, biological or botanical products, CC components, emollients, surfactants, thickening agents, silicone CC components, organic sunscreens, preservatives, neutralising agents, cC perfumes and pigments. (I) is useful as a hair care composition such as Sampoo, conditioner, anti-dandruff treatment, styling aids, styling conditioner, hair repair or treatment, serum, lotion, cream, pomade, or CC chemical treatment such as hair dye; skin care composition such as CC moisturising body wash, body wash, antimicrobial cleanser, skin protectant cream, body lotion, facial cream, moisturising cream, facial cleansers anti-acne treatment, facial toner, exfoliating cream, facial exfoliating gel, anti-acne treatment, facial toner, exfoliating cream, facial mask, after shave balm or sunscreen, skin care composition topically applied cover-the-counter drugs comprising anti-fungal treatments, anti-acne treatments, skin protectants, and antiperspirants; cosmetic composition composition composition makes treatments, skin protectants, and antiperspirants; cosmetic composition composition composition composition consectic composition composition consectic composition composition chosen from eye gel, high-melting point
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(CUEV/) CUEVAS W A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO
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CC lipstick, lipstick, lip gloss, lip balm, mascara, eyeliner, pressed CC powder formulation and foundation; nail care composition such as nail CC enamel, cuticle treatment, nail polish, nail treatment, or polish remover CC; an oral care composition such as toothpaste, mouth rinse, breath CC freshener, or whitening treatment; and over-the-counter pharmaceutical CC composition. The cosmetic composition is a mascara such as non-waterproof mascara, waterproof mascara, volumising mascara such as non-waterproof cc unling mascara, anhydrous waterproof mascara, water-based mascara, or cyelash or eyebrow treatment; a pressed powder; foundation such as water-cin-oil foundation, water-in-silicone foundation, oil-in-water foundation, canhydrous makeup stick, or cream-to-powder foundation. (I) has desired coharacteristics such as transparent film formation, hydrogel formation, comperties such as transparent film formation, hydrogel formation, comperties wiscoelastic behaviour, glass transition temperature, cloud temperature and decomposition temperature), and does not have any chemical modifications of the protein. This is the amino acid sequence of silk-elastin polymer SELP 47-E13 that may be used as the repeat sequence
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Best Local Similarity
Matches 752; Conserv
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GVGVPGKGVPGVG-PGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPG
                                            PGKGVPGVG-PGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVP
                                                                                                                                      VPGEGVPGVGVPGVGEPGVGVPGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVPGVGV
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90.6%;
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Pred. No. 6.9e-217;
0; Mismatches 26;
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                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                    products such as moisturizers, toners and makeup and nail care products such as polishes and polish removers. The invention enables controlled release of active agents. The present sequence is that of a protein which is related to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention relates to a novel system for providing controlled release delivery of an active agent which comprises a repeat sequence protein polymer and at least one active agent. The invention may be useful in a hair care composition, skin care composition, nail care composition, composition, oral care composition or over-the-counter pharmaceutical composition. The system may be useful in shampoos, gels, mousses and other hair care products, rinse-off conditioners, skin care mousses and other hair care products.
                                                                                                                                                                                                                                                                                                Sequence 884 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     System for providing controlled release delivery of active agent useful for incorporating active agents into personal care product compositions
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(MAZE/) MAZEAUD I.
                                                                                                                                                                                                         Local
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                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        g active agents into personal care product sequence protein polymer and active agent.
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Pred. No. 6.9e-217;
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RESULT 15
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ID ADU81666
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XX Prote Silk/
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14-MAY-2004; 2004US-00845936
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Chimeric.
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                                                                                                             Silk/elastin
                               US2004234609-A1.
                                                                                             Protein
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                25-NOV-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            peptide repeat sequence,
                                                                                                                            (first entry)
                                                                                                                                                           protein;
                                                                                            protein stabilization; cosmetics; textile; filter;
                                                                                                             SEQ ID NO:25
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                                                                                                                                                                                                                                                                                                                                                 CC sequence protein polymer (i.e., an engineered protein) and at least one cative agent, such as slicones, fragrances, dyes, tints, UV actives, CC sunscreens, lanolin, vitamins, bleaches, thickening agents, into the control polymer preferably comprises a repeating amino acid sequence comprised from elastin, collagen, abductin, byssus, flagelliform silk, CC derived from elastin, collagen, abductin, byssus, flagelliform silk, CC dragline silk, gluten high molecular weight subunit, titin, fibronectin, CC leminin, gliadin, glue polypeptide, ice nucleating protein, keratin, CC dragline silk, gluten high molecular weight subunit, titin, fibronectin, CC mucin, RNA polymerase II, resilin or mixtures thereof. The repeat sequence protein polymers provide a scaffolding to which active agents comprised a stative sequence protein sathing the protein sathing to which active agents as they overcome many of the CC hydrophobicity, solubility, and billow the use of polypeptide active agents in applications for which a total protein environment is desirable. The CC in applications for which a total protein environment is desirable. The CC in applications and a polypeptide active agents biomolecular conjugate by recompositions and a polypeptide active agent; biomoternials adapted for at least one predetermined desirable function which comprise one or more at least one predetermined desirable function which comprise one or more at least one predetermined desirable function which comprise one or more biomolecular conjugate by protein polymer and a polypeptide active agent; and a method of producing such biomaterials. The biomolecular conjugate are useful in compositions, and compositions such as over-the counter pharmaceutical corporate and compositions used for hair care, skin care, nail care, compositions, and compositions used for hair care, skin care, nail care, compositions, and compositions used for hair care, skin care, nail care, compositions are compositions and care compositions. The present sequence is given the sequence 
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Search completed: May 10, 2006, 10:39:41 Job time: 195 secs

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it is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1/iaa/5_COMB.pep:*

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3: /cgn2_6/ptodata/1/iaa/H_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*

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 Maximum Match 100%
Listing first 45 summaries
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GenCore version 5.1.8 (c) 1993 - 2006 Biocceleration Ltd.
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              US-08-806-029-19
US-08-337-641-18
US-08-337-631A-68
US-08-435-641-206-15
US-08-642-246-15
US-08-642-246-15
US-08-642-246-15
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US-08-177-5098-82
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US-08-444-791A-3
US-08-806-029-27
PCT-US95-02772-4
US-08-806-029-23
US-08-175-155-39
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US-08-482-0858-74
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US-08-482-088-74
US-08-482-088-74
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ALIGNMENTS	US-08-806-029-36	US-08-806-029-25	PCT-US96-06229-25	US-09-451-206-25	US-08-642-246-25	US-08-707-237A-103	PCT-US95-02772-6	US-08-806-029-29	US-08-212-237-6	US-08-707-237A-51	US-08-175-155-45	US-08-482-085B-80	US-08-477-509B-80	US-09-444-791A-80	US-09-444-791A-81	US-08-482-085B-81	US-08-707-237A-52
sequence IV8, App	36,	Sequence 25, Appl	Sequence 25, Appl	Sequence 25, Appl	Sequence 25, Appl	Sequence 103, App	Sequence 6, Appli	Sequence 29, Appl	Sequence 6, Appli	Sequence 51, Appl		Sequence 80, Appl	Sequence 80, Appl	Sequence 80, Appl	Sequence 81, Appl	Sequence 81, Appl	Sequence 52, Appl

RESULT 1

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US-08-806-029-19
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 Query Match
Best Local Similarity
Matches 765; Conserv
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                                                                                                                                                    APPLICATION NUMBER: US/08/806,029
FILING DATE: 24-FEB-1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/212,237
FILING DATE: 11-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Trecartin, Richard F.
REGISTRATION NUMBER: 31,801
                                                                                                                                                                                                                                                                                                                                                                                               ZIP: "94111
COMPUTER READABLE FORM:
COMPUTER TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
CURRENT APPLICATION UMBER: US/08/806,029
                                                                                     MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hohbach, Test,
STREET: Four Embarcadero Center,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Cappello, Joseph
APPLICANT: Stedronsky, Erwin R.
TITLE OF INVENTION: Synthetic Proteins for in vivo Drug
TITLE OF INVENTION: Delivery and Tissue Augmentation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: C
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                                                                                                      TOPOLOGY:
                                                                                                                                      TYPE: amino acid
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CITY: San Francisco
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   Conservative
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92.2%;
Score 3526; DB 2;
Pred. No. 5.6e-254;
0; Mismatches 13;
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Suite 3400
                                  Length 889;
   Indels
   52;
   Gaps
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| Sequence 68, Application US/08397633A |
| Patent No. 5773577 |
| Patent No. 5773577 |
| GENERAL INFORMATION: APPLICANT: Cappello, Joseph |
| TITLE OF INVENTION: PRODUCTS COMPRISING SUBSTRATESCAPABLE |
| TITLE OF INVENTION: OF ENZYMATIC CROSS-LINKING |
| NUMBER OF SEQUENCES: 105 |
| CORRESPONDENCE ADDRESS: CORRESPONDENCE ADDRESS: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT |
| STREET: San Francisco |
| STATE: California |
| COLUMBY: California |
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US-08-397-633A-68
COMPUTER READABLE FORM:
                         COUNTRY: USA
ZIP: 94111-4187
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; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-397-633A-68
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Matches 752
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LENGTH: 884 amino acids
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ATTORNEY/AGENT INFORMATION:
NAME: Rowland, Bertram I
REGISTRATION NUMBER: 20,015
REFERENCE/DOCKET NUMBER: A-58
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEPAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYDE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/397,633A
FILING DATE:
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Pred. No. 3.3e-248;
"" matches 26;
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Best Local :
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PALENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/435,641
FILING DATE: 05-MAY-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15, Application Patent No. 5817303
                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TOPOLOGY: lin
MOLECULE TYPE: |
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                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Rowland, Bertram I.
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A-
                                                                                                                                                                                                                                    Local Similarity
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CITY: San Francisco
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TVPE: amino acid
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Four Embarcadero Center, Suite 3400
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VENTION: Tissue Adhesive Using
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Pred. No. 3.3e-248;
0; Mismatches 26;
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US-08-707-237A-96
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FILING DATE: 03-SEP-19
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US
APPLICATION APPLICATION OF THE PRIOR DATE: 29-DEC-19
                                                                                                                                                                                                                                                                      APPLICANT: Ferrari, Franco A.
APPLICANT: Capello, Joseph
APPLICANT: Crissman, John W.
APPLICANT: Dorman, Mary A.
TITLE OF INVENTION: METHODS FOR PREPARING SYNTHETIC
TITLE OF INVENTION: REPETITIVE DNA
                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: BM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                    NUMBER OF SEQUENCES: 108
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hohbach, Test,
STREET: Four Embarcadero Center,
CITY: San Francisco
                                                                                                                                                                            STATE: C
                                                                   APPLICATION NUMBER:
                                                                                                                                                              COUNTRY: United States ZIP: 94111-4187
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  29-DEC-1993
                                                    03-SEP-1996
                                                                    US/08/707,237A
               US 08/175,155
                                                                                              Version #1.30
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Best Local (
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ETILING DATE: 29-OCT-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 06/927,258
FILING DATE: 04-NOV-1986
ATTORNEY/AGENT INFORMATION:
NAME: Trecartin, Richard F.
REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: A-55186-10,
TELEPHONE: (415) 781-1989
TELEPHONE: (415) 781-1989
TELEPAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: 910 277299
INFORMATION FOR SEQ ID NO:
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APPLICATION NUMBER: US
FILING DATE: 22-APR-1:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 06-NOV-1:
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MOLECULE TYPE:
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Similarity 90.6%;
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Pred. No. 3.3e-248;
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RESULT 5
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US-08-642-246-15
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                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: STEDRONSKY, Erwin R.
APPLICANT: CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive
TITLE OF INVENTION: Crosslinking
                                                                                                                                                                                                                     TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/642,246
                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: A6
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST,
STREET: Four Embarcadero Center,
                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: ROWLAND, Bertram I
REGISTRATION NUMBER: 20015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: CA
COUNTRY: US
ZIP: 94111
                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Four Embarca
CITY: San Francisco
                                                                                                           Local Similarity
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884 amino acids
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Pred. No. 3.3e-248;
0; Mismatches 26;
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RESULT 6
US-09-451-206-15
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        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEER, HOHBACH, TEST,
STREET: Four Embarcadero Center,
CITY: San Francisco
STATE: CA
COUNTRY: US
ZIP: 94111
                                                                                                                                         APPLICANT: STEDRONSKY, Erwin R.
CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive
 CURRENT APPLICATION DATA:
                                                                                                                       NUMBER OF SEQUENCES: 35
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           Version
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Suite 200
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MOLECULE TYPE: peptide
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SEQUENCE DESCRIPTION: S
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INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
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Local > 752;
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PRIOR APPLICATION DATA:
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                                                                          PGKGVPGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVP
                                                                 GVG-PGVG-PGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVG-PG
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                                                                                                                                                                                                                                                                                                                                LENGTH: 884 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A61127-1/BIR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/451,206
FILING DATE: 29-No. 6423333-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: ROWLAND, Bertram
                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                     82.6%;
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Pred. No. 3.3e-248;
0; Mismatches 26;
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PCT-US96-06229-15
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                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: peptide PCT-US96-06229-15
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GENERAL INFORMATION:
APPLICANT: STEDRONSKY, Erwin R.
APPLICANT: CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive Using Synthetic
TITLE OF INVENTION: Crosslinking
                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 752; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY ROWLAND, Bertram I
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A611
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: PCT/US96/06229
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CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT STREET: Four Embarcadero Center, Suite 200
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Conservative
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Pred. No. 3.3e-248;
0; Mismatches 26;
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US-08-175-155-47
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           APPLICATION NUMBER: US/08/
FILING DATE: 29-DEC-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: ROwland, Bertram I.
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A
                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                      APPLICANT: Ferrari, Franco A.
APPLICANT: Cappello, Joseph
APPLICANT: Crissman, John W.
APPLICANT: Dorman, Mary A.
ATITLE OF INVENTION: Methods for Preparing
TITLE OF INVENTION: Repetitive DNA
NUMBER OF SEQUENCES: 69
TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                              STREET: Four Embarcac
CITY: San Francisco
STATE: CA
COUNTRY: US
ZIP: 94111
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TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
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Best Local Similarity 67.4%;
Matches 755; Conservative
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TELEPAX: 415-398-3249
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 2557 amino acids
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Pred. No. 1.3e-236;
0; Mismatches 25;
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US-SOUTH US-	
-GR-477-509B-82 -GR-477-509B-82 sequence 82, Application US/08477509B sepulcann: Cappello, Joseph APPLICANN: Corpent, John APPLICANN: Corpent, John APPLICANN: Corpent, Mary A APPLICANN: Corpent, Mary A APPLICANN: Comman, John APPLICANN: Comman, Mary A APPLICANN: Comman, Mary A APPLICANN: Comman, Mary A APPLICANN: Compatible COMPUTER EADDALE FORM: MEDIUM TYPE: DEPPY disk COMPUTER SEADABLE FORM: MEDIUM TYPE: DAS/NS-DOS SOFTMARE: LAM PC-COMPATIBLE COMPUTER APPLICATION NOBER: US/08/175,155 SECONOMINER IDM COMPATIBLE COMPUTER APPLICATION NOBER: US/08/477,509B FILING DATE: 07-JUN-1995 SECONOMINER SECONOMINERS: US/08/477,509B FILING DATE: 03-DEC-1993 APPLICATION NUMBER: US/08/477,509B FILING DATE: 03-DEC-1993 APPLICATION NUMBER: US/08/477,509B FILING DATE: 03-DEC-1993 APPLICATION NUMBER: US/08/477,509B FILING DATE: 03-DEC-1993 BRIGG APPLICATION NUMBER: US/08/477,509B FILING DATE: 03-DEC-1993 APPLICATION NUMBER: US/08/477,509B FILING	

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67.4%;
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Pred. No. 1.3e-236;
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                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 07/609,716
FILING DATE: 06-NOV-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/269,429
FILING DATE: 09-NOV-1988
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/114,618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 03-SEP-1996
CLASSIFICATION: 435
FRIOR APPLICATION DATA: US 08/175,155
APPLICATION NUMBER: US 08/175,155
FILING DATE: 29-DEC-1993
PRIOR APPLICATION DATA: US 08/053,049
APPLICATION NUMBER: US 08/053,049
                                                                                 TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 2257 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Capello, Joseph
APPLICANT: Crissman, John W.
APPLICANT: Dorman, Mary A.
APPLICANT: Dorman, Mary A.
TITLE OF INVENTION: METHODS FOR PREPARING SYNTHETIC
TITLE OF INVENTION: REPETITIVE DNA
NUMBER OF SEQUENCES: 108
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                     FILING DATE: 04.NOV-1986
ATTORNEY/AGENT INFORMATION:
NAME: Trecartin, Richard F.
REGISTRATION NUMBER: 31,801
MOLECULE TYPE:
                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: A-TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 22-APR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/609,716
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APPLICATION NUMBER: US,
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CITY: San Francisco
STATE: California
                                        TYPE: amino acid
STRANDEDNESS: uni
                      TOPOLOGY:
                                                                                                                                                                         TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 94111-4187
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Four Embarcadero Center, Suite 3400
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-GPGAGAGSGAGAGSGAGAGSGVGVPGVGVPGV 744
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RESULT 11
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                                                                   Query Match
Best Local
                                                       Matches
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APPLICATION NUMBER: US 06/927,258
FILING DATE: 04-NOV-1986
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/114,618
FILING DATE: 29-OCT-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/053,049
FILING DATE: 22-APR-1993
PRIOR APPLICATION NUMBER: US 08/175,155
APPLICATION NUMBER: US 08/175,155
APPLICATION NUMBER: US 08/175,155
APPLICATION NUMBER: US 08/175,155
APPLICATION NUMBER: US 08/175,155
                                                                                                                                                                                                                             TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TOPOLOGY: line
MOLECULE TYPE: ]
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APPLICANT: Cappello, Joseph
APPLICANT: Crissman, John W.
TITLE OF INVENTION: No. 6018030el Peptides Comprising Repetitive
TITLE OF INVENTION: Units of Amino Acids and DNA Sequences Encoding
NUMBER OF SEQUENCES: 112
CORRESPONDENCE S. 112
CORRESPONDENCE S. 112
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APPLICANT:
                                                                                                                                                                                                                                                                           NAME: Trecartin, Richard F.
REGISTRATION NUMBER: 31.801
REFERENCE/DOCKET NUMBER: A-55186-6/RFT/MTK
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
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APPLICATION NUMBER: US,
FILING DATE: 07-JUN-19:
CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 29-DEC-1993
ATTORNEY/AGENT INFORMATION:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
                                                                                                                                                                                           TYPE: amino acid
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                                                                                                                                                                                                             LENGTH:
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Chambers, James
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Pred. No. 1.3e-236;
0; Mismatches 25;
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Suite 3400
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                                                                                                                                           Query Match
Best Local Similarity
Matches 755; Conserv
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INFORMATION FOR SEQ ID NO: 82:
SEQUENCE CHARACTERISTICS:
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FILING DATE: 07-UW-1995
APPLICATION NUMBER: US 08/175,155
FILING DATE: 29-DEC-1993
APPLICATION NUMBER: US 08/053,049
FILING DATE: 22-DRR-1993
APPLICATION NUMBER: US 07/114,618
FILING DATE: 22-CCT-1987
APPLICATION NUMBER: US 06/927,258
FILING DATE: 04-NOV-1986
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Trecartin, Richard F.
REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: A-55186-11/RFT/BTC
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/09/444,791A
FILING DATE: 22-No. 6355776-1999
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr Hohbach Test Albritton & Herbert
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GAGSGAGAGSGAGAGSGAGAGSGAGAGSGAGAGSGVPGVPGVPGVPGVPGVPGVP
                                                                       STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
COUNTRY: US
                                                                                                                                                                                                                                                                        STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                            TYPE: amino acid
                                                                                                                                                                                                                                                                                                                             LENGTH: 2257 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cappello, Joseph
Crissman, John W.
OF INVENTION: No. 6355776el Peptides Comprising Repetitive
Units of Amino Acids and DNA Sequences Encoding the
                                                                                                                                               Conservative
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Pollock, Thomas J.
                  79.0%;
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                                                                                                                                           Score 3296; DB 2;
Pred. No. 1.3e-236;
0; Mismatches 25;
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RESULT 13
US-08-397-633A-54
; Sequence 54, Application
; Patent No. 5773577
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US-08-397-633A-54
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Best Local Similarity
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ZOTE: 94111-4187

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vi
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: 910 277299
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: ROWLAND, Bertram I
REGISTRATION NUMBER: 20,015
REFERENCE/DOCKET NUMBER: A-58848-1/BIR PROP-011-1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 877 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Cappello, Joseph
TITLE OF INVENTION: PRODUCTS COMPRISING SUBSTRATESCAPABLE
TITLE OF INVENTION: OF ENZYMATIC CROSS-LINKING
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CITY: San Francisco
STATE: California
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TELEFAX: (415) 398-3249
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
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              VG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVVPGVGVPGKGVPGV
                                                        VGVPGVGVPGVGVPGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVPGVPGVPGVPGVPGV
                                             E: FLEHR, HOHBACH, TEST, ALBRITTON 4 Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                      Conservative
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89.9%;
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    Mismatches

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Pred. No. 4.5e-236;
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                                                                                        ; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-212-237-4
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US-08-212-237-4
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                                              Matches
                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                              ZIP: 94111-4187

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, V6
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Cappel
                                                                                                                                                                     TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                            NAME: Rowland, Bertram I REGISTRATION NUMBER: 20,015 REFERENCE/DOCKET NUMBER: A-1TLECOMMUNICATION INFORMATION: TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/212,237
FILING DATE: 11-MAR-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Cappello, TITLE OF INVENTION:
                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                LENGTH:
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                                            78.0%; ilarity 88.9%; Conservative
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Synthetic Proteins As
                                                                                                                                                                                                                     A-58847/BIR
                                            0
                                          Score 3253; DB 1; Length 8 Pred. No. 8.5e-234; 0; Mismatches 13; Indels
                                                                                                                                                                                                                                                                                                               Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Implantables
                                                                 Length 832;
                                             80;
                                            Gaps
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                                                                                                                                                                                             RESULT 15
US-08-806-029-27
 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                STREET: Four Embarcade:
CITY: San Francisco
STATE: California
COUNTRY: United States
                                                                             ADDRESSEE:
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Four Embarcadero Ce
Floppy disk
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Sequence 27, Application US/08806029

Sequence 27, Application US/08806029

Patent No. 6380154

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Stedronsky, Erwin R.
TITLE OF INVENTION: Synthetic Proteins for in vivo Drug
TITLE OF INVENTION: Delivery and Tissue Augmentation
NUMBER OF SEQUENCES: 36

CORRESPONDENCE ADDRESS:
ANDRESSETT. FIRST WORKERS ADDRESS:
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NAME: Trecartin, Richard F.
REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: A-58847-2/RFT/MTK
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEPAX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 832 amino acids
TYPE: amino acids
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Best Local Similarity 88.9%;
Matches 742; Conservative
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TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: protein
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APPLICATION NUMBER: US 08/212,237
FILING DATE: 11-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Trecartin, Richard F.
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRINT APPLICATION DATA:
APPLICATION NUMBER: US/08/806,029
FILING DATE: 24-FEB-1997
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Pred. No. 8.5e-234;
0; Mismatches 13; Indels 8
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Search completed: May 10, 2006, 10:45:25 Job time : 55 secs

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Result
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Match
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

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Listing first 45 summaries
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 GenCore version 5.1.8 (c) 1993 - 2006 Biocceleration Ltd.
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US-10-441-965-19
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US-08-966-029-19
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Sequence 19, Application US/10800179

Publication No. US20040180027A1

GENERAL INFORMATION:
APPLICANT: Cuevas, William A.
APPLICANT: Cuevas, William A.
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj

TITLE OF INVENTION: Use of Repeat Sequence Protein Polymers in Personal Care

TITLE OF INVENTION: Compositions

FILE REFERENCE: DOC 0057 PA / GC792-4 / DC 5074

CURRENT APPLICATION NUMBER: US/10/800,179

CURRENT FILING DATE: 2004-03-12

PRIOR APPLICATION NUMBER: 60/454,077

PRIOR APPLICATION NUMBER: 60/454,077

PRIOR FILING DATE: 2003-03-12

NUMBER OF SEQ ID NO 19

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ORGANISM: Unknown
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100.0%; Pred. No. 6.6e-248;
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APPLICANT: Kumar, Manoj
APPLICANT: Kumar, Manoj
APPLICANT: Kumar, Manoj
APPLICANT: Mazeaud, Isabelle
APPLICANT: Mazeaud, Isabelle
APPLICANT: Christiano, Steven P.
TITLE OF INVENTION: Controlled Release of Active Agents Utilizing Repeat Sequence
TITLE OF INVENTION: Protein Polymers
FILE REFERENCE: DOC 0077 PA/GG 792-6/DC 5110
CURRENT APPLICATION NUMBER: US/10/845,775A
CURRENT FILING DATE: 2004-05-14
PRIOR APPLICATION UNMBER: US 60/470,465
PRIOR FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 31
SOFTWARE: Patentin version 3.2
SEQ ID NO 19
LENGTH: 780
                                                                                                  Query Match
Best Local Similarity
Matches 780; Conserv
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                                                                                                  100.0%; Score 4173; DB 5;
100.0%; Pred. No. 6.6e-248;
tive 0; Mismatches 0;
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APPLICANT: Collier, Katherine D.
APPLICANT: Collier, Katherine D.
APPLICANT: Collier, Katherine A.
APPLICANT: Collier, Katherine A.
APPLICANT: Kumar, Manoj A.
APPLICANT: Kumar, Manoj A.
TITLE OF INVENTION: Repeat Sequence Protein Polymer Active Agent Conjugates, Meth
TITLE OF INVENTION: and Uses
FILE REFERENCE: DOC0068PA/DC5058/GC792
CURRENT APPLICATION NUMBER: US/10/845,936A
CURRENT FILING DATE: 2004-05-14
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 37
SOFTWARE: PatentIn version 3.2
SEQ ID NO 19
LENGTH: 780
TYPE: PRT
ORGANISM: unknown
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US-10-845-936A-19
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Pred. No. 6.6e-248;
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US-10-939-036-19

Sequence 19, Application US/10939036

Publication No. US20050142094A1

GENERAL INFORMATION:

APPLICANT: Kumar, Manoj

TITLE OF INVENTION: Use of Repeat Sequence Protein Polititle OF INVENTION: Use of Repeat Sequence Protein Polititle REFERENCE: GC792-4-2

CURRENT APPLICATION UMBER: US/10/939,036

CURRENT APPLICATION NUMBER: US/10/939,036

CURRENT APPLICATION NUMBER: US/10/939,036

PRIOR APPLICATION NUMBER: US/0800,179

PRIOR FILING DATE: 2004-03-12

PRIOR APPLICATION NUMBER: US/0800,179

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RESULT 6
US-08-806-029-19
; Sequence 19, Application US/08806029
; Publication No. US20020045567A1
; GENERAL INFORMATION:
APPLICANT: Cappello, Joseph
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SEQ ID NO 19
LENGTH: 780
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TYPE: PRT
ORGANISM: Unknown
     APPLICANT:
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Stedronsky, Erwin R.
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TYPE: amino acid
STRANDEDNESS: unkno
TOPOLOGY: unknown
MOLECULE TYPE: protei
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REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: A-58847-2/RFT/MTK
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEPAX: (415) 388-3249
INFORMATION FOR SEQ ID NO: 19:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: FORM:
COMPUTER: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
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CITY: San Francisco
STATE: California
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TITLE OF INVENTION: Delivery and Tissue Augmentation
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
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APPLICATION NUMBER: US 0
FILING DATE: 11-MAR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 889 amino acids
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Pred. No. 2.6e-208;
0; Mismatches 13;
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RESULT 7
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US-10-117-931-15
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Publication No. US20030104589A1
GENERAL INFORMATION:
  Query Match
                                                                                                               TELEFAX: 415-398-3249 INFORMATION FOR SEQ ID NO: 15: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.
CURRENT APPLICATION NATA:
APPLICATION NUMBER: US/10/117,931
FILING DATE: 05-Apr-2002
CLASSIFICATION: «Unknown»
                                                                                                                                                  ATTORNEY/AGENT INFORMATION:

NAME: ROWLAND, BETTEAM I

REGISTRATION NUMBER: 20015

REFERENCE/DOCKET NUMBER: A61127-1/BIR

TELECOMMUNICATION INFORMATION:

TELEPHONE: 415-781-1989
                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/642,246
FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST,
STREET: Four Embarcadero Center,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: STEDRONSKY, Erwin R.
CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive Using Synthetic
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                                                    MOLECULE TYPE:
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TYPE: amino acid
STRANDEDNESS: single
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                                                              TOPOLOGY: linear
                                       TYPE: peptide DESCRIPTION: :
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US-10-800-179-25
; Sequence 25, Application US/10800179
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                                                                                                    GENERAL INFORMATION:
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj
ITILE OF INVENTION: Use of Repeat Sequence Protein
TITLE OF INVENTION: Compositions
FILE REFERENCE: DOC 0057 PA / GC792-4 / DC 5074
CURRENT APPLICATION NUMBER: US/10/800,179
CURRENT FILING DATE: 2004-03-12
PRIOR APPLICATION NUMBER: 60/454,077
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Conservative
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smatches 26;
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NUMBER OF SEQ ID NOS: 31
SOFTWARE: PatentIn version 3.2
SEQ ID NO 25
LENGTH: 884
TYPE: DEF
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Best Local (
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ORGANISM: Unknown
FEATURE:
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Similarity 90.6%;
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Pred. No. 1.5e-203;
0; Mismatches 26;
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RESULT

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Sequence 25, Application US/10845775A

Publication No. US20040228913A1

GENERAL INFORMATION:
APPLICANT: Kumar, Manoj

APPLICANT: Mazeaud, Isabelle
APPLICANT: Mazeaud, Isabelle
APPLICANT: Christiano, Steven P.
TITLE OF INVENTION: Controlled Release of Active Age
TITLE OF INVENTION: Protein Polymers

FILE REFERENCE: DOC 0077 PA/GC 792-6/DC 5110

CURRENT APPLICATION NUMBER: US/10/845,775A

CURRENT FILING DATE: 2004-05-14

PRIOR APPLICATION NUMBER: US 60/470,465

PRIOR APPLICATION SUMBER: US 60/470,465

PRIOR APPLICATION SUMBER: US 60/470,465

PRIOR FILING DATE: 2003-05-14

NUMBER OF SEQ ID NOS: 31

SOFTWARE: Patentin version 3.2

SEQ ID NO 25

LENGTH: 884
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; ORGANISM: unknown
; FEATURE:
; OTHER INFORMATION: artificial glycine-rich peptide repeat sequence
US-10-845-775A-25
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Best Local Similarity
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Pred. No. 1.5e-203;
0; Mismatches 26;
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Sequence 25, Application US/10845936A
Publication No. US20040234609A1
GENERAL INFORMATION:
APPLICANT: COllier, Katherine D.
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj A.
TITLE OF INVENTION: Repeat Sequence Protein Pol
TITLE OF INVENTION: and Uses
CURRENT APPLICATION UNMBER: US/10/845,936A
CURRENT FILING DATE: 2004-05-14
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR FILING DATE: 2003-05-14
ORGANISM: US/10/845,936A
CURRENT FILING DATE: 2003-05-14
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR FILING DATE: 2003-05-14
CURBER OF SEQ ID NOS: 37
SOFTWARE: PatentIn version 3.2
SEQ ID NO 25
LENGTH: 884
TYPE: PRT
CORGANISM: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 10
US-10-845-936A-25
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US-10-845-936A-25
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Best Local S
Matches 752
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Local Similarity 90.6%;
hes 752; Conservative
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                                                                                           GVPGVGVPGVGVPGVGVPGBGVPGVGVPGVGBPGVGVPGAGAGSGAGAGS
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Pred. No. 1.5e-203;
0; Mismatches 26;
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CURRENT APPLICATION NUMBER: US/10/939,036
CURRENT FILING DATE: 2004-09-10
PRIOR APPLICATION NUMBER: US 10/800,179
PRIOR FILING DATE: 2004-03-12
PRIOR FILING DATE: 2004-03-12
PRIOR APPLICATION NUMBER: US 60/454,077
PRIOR APPLICATION NUMBER: US 60/454,077
PRIOR APPLICATION NUMBER: US 60/454,077
PRIOR FILING DATE: 2003-03-12
NUMBER OF SEQ ID NOS: 37
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WS-10-939-036-25
(Sequence 25, Application US/10939036
; Publication No. US20050142094A1
; GENERAL INFORMATION:
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Best Local
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ORGANISM: Unknown
FEATURE:
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Local Similarity 90.6%;
es 752; Conservative
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Pred. No. 1.5e-203;
0; Mismatches 26;
                                                                                                                                                                                                                                                                                                            DB 5; Length 884;
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; OTHER INFORMATION: ; OTHER INFORMATION: US-10-845-936A-37
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US-10-845-936A-37
                                                                                                                                                                                                             APPLICANT: Collier, Katherine D.
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj A.
TITLE OF INVENTION: Repeat Sequence Protein Polymer Active Agent Conjugates, ITILE OF INVENTION: and Uses
FILE REFERENCE: DOC0068PA/DC5058/GC792
CURRENT APPLICATION NUMBER: US/10/845,936A
CURRENT FILING DATE: 2004-05-14
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR FILING DATE: 2003-05-14
RUMBER OF SEQ ID NOS: 37
SOPTWARE: PatentIn version 3.2
SEQ ID NO 37
LENGTH: 983
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 37, Application US/10845936A Publication No. US20040234609A1 GENERAL INFORMATION:
                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                          FEATURE:
                                                                                                                                                                                       ORGANISM: unknown
                                                                                                                                                                                                      TYPE: PRT
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                                         GSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVG-PGVG-PGVG-PGAGAGS 114
                           VGVPGVGVPGVGVPGVGVPGVGVPGVGEPGVGVPGAGAGSGAGAGSGAGAGSGAGAGSGV
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                                                                                 Conservative
                                                                                                                                              CBPhexamer-SELP47K: silk, elastin protein polymer sequence
                                                                                            90.6%;
                                                                            Score 3448; DB 5;
Pred. No. 1.6e-203;
0; Mismatches 26;
                                                                              Indels
                                                                                                        Length
                                                                                                                                                            and cellulose binding peptide
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                                  Sequence 32, Application US/10845936A
Publication No. US20040234609A1
GENERAL INFORMATION:
APPLICANT: COllier, Katherine D.
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj A.
TITLE OF INVENTION: Repeat Sequence Protein Pol
TITLE OF INVENTION: and Uses
FILE REFERENCE: DOC0068PA/DC5058/GC792
CURRENT APPLICATION NUMBER: US/10/845,936A
CURRENT FILING DATE: 2004-05-14
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR FILLING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 37
SOFTWARE: Patentin version 3.2
SEQ ID NO 32
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LENGTH: 1027
TYPE: PRT
ORGANISM: unknown
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Methods

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RESULT 14

US-10-845-936A-33

Sequence 33, Application US/10845936A

; Publication No. US20040234609A1

; GENERAL INFORMATION:

; APPLICANT: Collier, Katherine D.

; APPLICANT: Cuevas, William A.

; APPLICANT: Kumar, Manoj A.
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TITLE OF INVENTION: Repeat Sequence Protein Pol
TITLE OF INVENTION: and Uses
FILE REFERENCE: DOCOGGBPA/DC5058/GC792
CURRENT APPLICATION NUMBER: US/10/845,936A
CURRENT FILING DATE: 2004-05-14
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 37
SOFTWARE: PatentIn version 3.2
SEQ ID NO 33
LENGTH: 1105
TYPE: PRT
ORGANIUM: unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Collier, Katherine D.
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj A.
TITLE OF INVENTION: and Uses
TITLE OF INVENTION: and Uses
TILE REFERENCE: DOC0068PA/DC5058/GC792
CURRENT APPLICATION NUMBER: US/10/845,936A
CURRENT APPLICATION NUMBER: 2004-05-14
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 37
COMPTUANDE: Datantin varsion 1 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: unknown
FEATURE:
OTHER INFORMATION: GFP-SELP47K: silk, elastin and green fluorescent protein peptides
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Search completed: May 10, Job time : 171 secs 2006, 10:48:14

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Database
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                 Published Applications AA New:*

1: /SIDSS/ptodata/2/pubpaa/US06 NEW_PUB.pep1:*

2: /SIDSS/ptodata/2/pubpaa/US06 NEW_PUB.pep:*

3: /SIDSS/ptodata/2/pubpaa/US07 NEW_PUB.pep:*

4: /SIDSS/ptodata/2/pubpaa/US08 NEW_PUB.pep:*

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9: /SIDSS/ptodata/2/pubpaa/US11 NEW_PUB.pep:*

10: /SIDSS/ptodata/2/pubpaa/US11 NEW_PUB.pep:*

11: /SIDSS/ptodata/2/pubpaa/US11 NEW_PUB.pep:*

12: /SIDSS/ptodata/2/pubpaa/US10 NEW_PUB.pep:*
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                               Minimum Match 0%
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Listing first 45 summaries
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US-11-053-100-30
US-11-053-100-30
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US-11-201-606-15
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	Sequence 48, Appl	•	Sequence 145, App	Sequence 135, App	•	Sequence 36, Appl	•	•	•	•	•	Sequence 16, Appl	57,	•	Sequence 34, Appl	Sequence 37, Appl	-	•	•	•	•	•	Sequence 45, Appl

ALIGNMENTS

Title:

Run on:

Sequence 15, Application US/11201606 Publication No. US20060029638A1 STATE: CA
COUNTRY: US
ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION UNWHER: US/11/201,606
FILING DATE: 2005-08-10
CLASSIFICATION
PRICA PAPLICATION DATA:
APPLICATION UNWHER: US/08/642,246
FILING DATE: 1996-05-02
CLASSIFICATION NUMBER: US/08/642,246
FILING DATE: 1996-05-02 TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS: CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
NAME: ROWLAND, Bertram I
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A611
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989 GENERAL INFORMATION: APPLICANT: STEDRONSKY, Erwin R.
APPLICANT: CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive
TITLE OF INVENTION: Crosslinking TYPE: amino ac STRANDEDNESS: TOPOLOGY: lin CORRESPONDENCE ADDRESS: MOLECULE NUMBER OF SEQUENCES: STREET: Four Embarca CITY: San Francisco LENGTH: ADDRESSEE: amino acid 884 amino acids E: FLEHR, HOHBACH, TEST, Four Embarcadero Center, linear single 15: A61127-1/BIR ALBRITTON Suite 200 Using Synthetic #1.30 & HERBERT

Result No.

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RESULT 2
US-11-201-606-25
; Sequence 25, Application US/11201606
; Publication No. US20060029638A1
; GENERAL INFORMATION:
; APPLICANT: STEDRONSKY, Erwin R.
; APPLICANT: CAPPELLO, Joseph
; TITLE OF INVENTION: Tissue Adhesive Using Syn
; TITLE OF INVENTION: Crosslinking
; NUMBER OF SEQUENCES: 35
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Pred. No. 6.6e-230;
0; Mismatches 26;
                  Synthetic
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Best Local Similarity
Matches 668; Conserv
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/642
FILING DATE: 1996-05-02
CLASSITICATION:
ATTORNEY/AGENT INFORMATION:
NAME: ROWLAND, BERTIAM I
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A611
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
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ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON
STREET: Four Embarcadero Center, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 1002 amino acids
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CITY: San Francisco
STATE: CA
COUNTRY: US
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STRANDEDNESS: sir
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415-398-3249
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RESULT 3
US-11-201-606-30
US-11-201-606-30
; Sequence 30, Application US/11201606
; Publication No. US20060029638A1
; GENERAL INFORMATION:
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                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC_DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/201,606
FILING DATE: 2005-08-10
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/642,246
FILING DATE: 1996-05-02
CLASSIFICATION:
APPLICATION NUMBER: US/08/642,246
FILING DATE: 1996-05-02
CLASSIFICATION NUMBER: 20015
REGISTRATION NUMBER: 20015
REGISTRATION NUMBER: A61127-1/BIR
REFERENCE,DOCKET NUMBER: A61127-1/BIR
                                                                                                                                                                                                 ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                               APPLICANT: STEDRONSKY, Erwin R.
APPLICANT: CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive
TITLE OF INVENTION: Crosslinking
                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
       TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                  CITY: San Fr
STATE: CA
COUNTRY: US
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Four Embarcadero Center,
415-398-3249
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Suite 200
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STRANDEDNESS: sir
TOPOLOGY: linear
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nes 642; Conserv
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73.9%;
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Pred. No. 4.1e-185;
4; Mismatches 93;
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US-11-201-606-34
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Publication No. US20060029638A1
GENERAL INFORMATION:
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Best Local Similarity 72.5%;
Matches 642; Conservative
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APPLICANT: CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive Using Sy.
TITLE OF INVENTION: Crosslinking
NUMBER OF SEQUENCES: 35
CORRESPONDENCE ADDRESS:
ADDRESSE: FOLERR, HOHBACH, TEST, ALBRITTON
STREET: FOUR Embarcadero Center, Suite 200
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: ROWLAND, Bertram I
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A61
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: US/
FILING DATE: 1996-05-02
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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VG-PGVG-PGAGAGSGAGAGSGAGA-GSGAGAGSGVGVPGVGVPGVGVPGKGVPGKGVPGVGPGV
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Pred. No. 2.5e-181;
4; Mismatches 104;
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Suite 200
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CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 09/812,382
PRIOR FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: US 60/190,659
PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
SOFTWARE: PatentIn version 3.3
SEQ ID NO 46
                                                                                                                 ; OTHER INFORMATION: Synthetic Construct
; FEATURE:
; NAME/KEY: MISC FEATURE
; LOCATION: (1).-(1186)
; OTHER INFORMATION: pET15b-SD3-ELP1-180-throm-Androgen receptor ligand
US-11-053-100-46
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Sequence 46, Application US/11053100
Publication No. US2005025554A1
GENERAL INFORMATION:
APPLICANT: CHILKOTI, Ashutosh
TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE
FILE REFERENCE: 4176-101 CIP
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Best Local S
Matches 542
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TYPE: PRT
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                                                  Score 2091.5; DB 11;
Pred. No. 8.4e-137;
2; Mismatches 223; I
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Sequence 53, Application US/11053100
Publication No. US2005025554A1
GENERAL INFORMATION:
APPLICANT: CHILKOTI, Ashutosh
TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE
FILE REFERENCE: 4176-101 CIP
CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILING DATE: 2005-02-08
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PRIOR FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: US 60
PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
SOFTWARE: PatentIn version 3.3
SEQ ID NO 53
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; OTHER INFORMATION: pET15b-SD1-ELP1-180-throm-G
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FEATURE:
NAME/KEY: MISC FEATURE
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ORGANISM: Artificial
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Pred. No. 1.7e-136;
3; Mismatches 223; I
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CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 09/812,382
PRIOR EILING DATE: 2001-03-20
PRIOR FILING DATE: 2000-03-20
PRIOR FILING DATE: 2000-03-20
PRIOR FILING DATE: 2000-03-20
PRIOR FILING DATE: 58
SOFTWARE: Patentin version 3.3
SEQ ID NO 26
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TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE
FILE REFERENCE: 4176-101 CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: MISC_FEATURE
LOCATION: (1)..(939)
OTHER INFORMATION: pET32a-SD15-ELP1-180-EK-Insulin
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ORGANISM: Artificial
FEATURE:
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                                                                       GAGSGVGVPGVGVPGVGVPGKGVPGVG-PGVG-PGVG-PGAGAGSGA------
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Pred. No. 1.7e-136;
2; Mismatches 225;
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FEATURE:
COTHER INFORMATION: Synthetic Construct
FEATURE:
NAME/KEY: MISC FEATURE
LOCATION: (1)...(1174)
COTHER INFORMATION: pET15b-SD5-ELP1-180-throm-Tobacco etch virus
US-11-053-100-43
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NUMBER OF SEQ ID NOS: 58
SOFTWARE: PatentIn version 3.3
SEQ ID NO 43
                                                         Query Match
Best Local Similarity
Matches 539; Conserv
                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: CHILKOTI, Ashutosh
TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE
FILE REFERENCE: 4176-101 CIP
CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILING DATE: 2005-02-08
                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 09/812,382 PRIOR FILING DATE: 2001-03-20
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                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Artificial
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Pred. No. 2e-136;
2; Mismatches 225;
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Sequence 50, Application US/11053100
Publication No. US2005025554A1
GENERAL INFORMATION:
APPLICANT: CHILKOTI, Ashutosh
TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE
FILE REFERENCE: 4176-101 CIP
CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 09/812,382
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PRIOR APPLICATION NUMBER: US 60
PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
SOFTWARE: PatentIn version 3.3
SEQ ID NO 50
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TYPE: PRT
ORGANISM: Artificial
FEATURE:
OTHER INFORMATION: Synthetic Co
FEATURE:
FEATURE:
NAME/KEY: MISC FEATURE
NAME/KEY: MISC TEATURE
TO THE TONE (1) ... (1225)
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Best Local Similarity
Matches 539; Conserv
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OTHER INFORMATION: pET15b-SD5-ELP1-180-throm-Estrogen receptor ligand binding domain
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Pred. No. 2.1e-136;
2; Mismatches 225;
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US-11-053-100-56
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CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 09/812,382
PRIOR FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: US 60/190,659
PRIOR FILING DATE: 2000-03-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 56, Application US/11053100
Publication No. US2005025554A1
GENERAL INFORMATION:
APPLICANT: CHILKOTI, Ashutosh
                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.3 SEQ ID NO 56
                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            IOCATION: (1)..(1329)
OTHER INFORMATION: perisb-sds-elp1-180-throm-1-Deoxy-D-Xylulose 5-Phosphate
OTHER INFORMATION: Reductoisomerase Peptide
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NAME/KEY: MISC FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Artificial
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Pred. No. 2.2e-136;
2; Mismatches 225;
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PRIOR FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: US 60/
PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
SOFTWARE: Patentin version 3.3
SEQ ID NO 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 11
US-11-053-100-32
                                                                                                                 US-11-053-100-32
                                                       Query Match
Best Local S
Matches 492
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publication No. US2005025554A1
GENERAL INFORMATION:
APPLICANT: CHILKOTI, Ashutosh
TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE
                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: 4176-101 CIP
CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 09/812,382
PRIOR FILING DATE: 2001-03-20
                                                                                                                     OTHER INFORMATION: Synthetic Construct FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (1)...(657)
OTHER INFORMATION: pET17b-ELP4-120-Throm-T20 peptide
                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial
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                                                                                                                                                                                                                                            LENGTH: 657
                                                                     Local Similarity
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                                                       Score 1913.5; DB 11;
Pred. No. 8.3e-125;
0; Mismatches 65; I
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APPLICANT: CHILKOTI, ASHUTOSH
FITTLE OF INVENTION: FUSION PEPTIDES ISOLATABLE BY
FILE REFERENCE: 4176-101 CIP
CURRENT APPLICATION UNMBER: US/11/053,100
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 09/812,382
PRIOR FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: US 60/190,659
PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
SOFTWARE: PatentIn version 3.3
SEQ ID NO 25
LENGTH: 639
TYPE: PRT
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US-11-053-100-25
                                                                                                                                   Sequence 25, Application US/11053100 Publication No. US20050255554A1 GENERAL INFORMATION:
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Best Local :
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OTHER INFORMATION: PET
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OTHER INFORMATION: Synthetic
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Pred. No. 1.3e-124;
0; Mismatches 66;
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Publication No. US20050255554A1

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PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
SOFTWARE: PatentIn version 3.3
SEQ ID NO 29
LENCTH: 654
TYPE: PRT
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Best Local :
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TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE BY PHASE TRANSITION
FILE REFERENCE: 4176-101 CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/11/053,100 CURRENT FILING DATE: 2005-02-08 PRIOR APPLICATION NUMBER: US 09/812,382 PRIOR FILING DATE: 2001-03-20
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              575 G-PGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGVGVPGVPGVGVPGVGVPGKG
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Similarity 59.7%;
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Pred. No. 1.3e-124;
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; OTHER INFORMATION: pET17b-ELP4-120-TEV(Q/Y)-T20 peptide
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US-11-053-100-38
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Publication No. US2005025554A1
GENERAL INFORMATION:
APPLICANT: CHILKOTI, Ashutosh
TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE BY PHASE TRANSITION
FILE REFERENCE: 4176-101 CTP
CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 09/812,382
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Best Local S
Matches 490
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NUMBER OF SEQ ID NOS: 58
SOFTWARE: PatentIn version 3.3
SEQ ID NO 38
LENGTH: 656
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ORGANISM: Artificial
FEATURE:
OTHER INFORMATION: Synthetic Construct
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Pred. No. 2.5e-124;
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Qy 123 Db 94	Qy 67 Db 49	Qy 11 Db 3	Query Match Best Local Si Matches 490;	; FEATURE: ; NAME/KEY: MI ; LOCATION: (1 ; OTHER INFORM US-11-053-100-35		SEQ ID NO 35 LENGTH: 657	NUMBER	PRIOR	CURREN	TITLE OF FILE REFE	Publicatio GENERAL IN	RESULT 15 US-11-053-100-35 ; Seguence 35, A		Оу 743	Db 514	Qy 687	Oy 631 Db 473	Db 432	Qy 575	Db 391	Оу 518	Db 350	Qy 461	Db 309	404
GAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVG-PGVG-PGVG-PGVG-P	GAGAGSGYGYPGYGYPGYGYPGKGYPGYG-PGYG-PGYG-PGYG-P	GSGVGVPGVGVPGVGVPGKGVPGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGS 66	/ Match 45.7%; Score 1906.5; DB 11; Length 657; Local Similarity 59.8%; Pred. No. 2.5e-124; 1es 490; Conservative 0; Mismatches 65; Indels 265; Gaps 67;	FEATURE: NAME/KEY: MISC_FEATURE LOCATION: (1)(657) OTHER INFORMATION: pET17b-ELP4-120-TEV(Q/S)-T20 peptide .1-053-100-35	ORGANISM: Artificial FEATURE: POTHER INFORMATION: Synthetic Construct	57	Q ID NOS: 58 tentIn version	FILING DATE: 2001-03-20 FILING DATE: 2000-03-20	T FILING DATE: 2005-02-08 APPLICATION NUMBER: US 09/812,382 STITUS DATE: 2005-02-08	ION: FUSION F 4176-101 CIE	NO. US20050255554A1 DRMATION:	-35 , Application US/11053100		GVGVPGKGVPGVG-PGVG-PGVG-PGAGAGSGAGA 778		PGKGVPGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVP 742	VPGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGV 686	ĠvÞĠŸĠvÞĠŸĠvBĠŸĠvPĠŸĠv	G-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVP	VGVPGVGVPGVGVPGVGVPGVGVPGVGVPGVGVPGVGV	VG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVP	GVGVPGVGVPGVGV	GVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVP	PGVGVPGVGV	

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Search completed: May 10, 2006, 10:50:15
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